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#### (54) Primers for synthesising full-length cDNA and their use

(57) Primers for synthesizing full-length cDNAs and their use are provided.

5602 cDNA encoding a human protein has been isolated and nucleotide sequences of 5'-, and 3' -ends of the cDNA have been determined. Furthermore, prim-

ers for synthesizing the full-length cDNA have been provided to clarify the function of the protein encoded by the cDNA. The full-length cDNA of the present invention containing the translation start site provides information useful for analyzing the functions of the protein.

#### Description

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#### FIELD OF THE INVENTION

[0001] The present invention relates to a polynucleotide encoding a novel protein, a protein encoded by the polynucleotide, and new uses of these.

#### BACKGROUND OF THE INVENTION

[0002] Currently, the sequencing projects, the determination and analysis of the genomic DNA of various living organisms have been in progress all over the world. The whole genomic sequences of more than 10 species of prokaryotes, a lower eukaryote, yeast, and a multicellular eukaryote, C. elegans are already determined. As to human genome, which is supposed to be composed of three thousand million base pairs, the world wide cooperative projects have been under way to analyze it, and the whole structure is predicted to be determined by the years 2002-2003. The aim of the determination of genomic sequence is to reveal the functions of all genes and their regulation and to understand living organisms as a network of interactions between genes, proteins, cells or individuals through deducing the information in a genome, which is a blueprint of the highly complicated living organisms. To understand living organisms by utilizing the genomic information from various species is not only important as an academie subject, but also socially significant from the viewpoint of industrial application.

[0003] However, determination of genomic sequences itself cannot identify the functions of all genes. For example, as for yeast, only the function of approximately half of the 6000 genes, which is predicted based on the genomic sequence, was able to be deduced. As for human, the number of the genes is predicted to be approximately one hundred thousand. Therefore, it is desirable to establish "a high throughput analysis system of the gene functions" which allows us to identify rapidly and efficiently the functions of vast amounts of the genes obtained by the genomic sequencing.

[0004] Many genes in the eukaryotic genome are split by introns into multiple exons. Thus, it is difficult to predict correctly the structure of encoded protein solely based on genomic information. In contrast, cDNA, which is produced from mRNA that lacks introns, encodes a protein as a single continuous amino acid sequence and allows us to identify the primary structure of the protein easily. In human cDNA research, to date, more than one million ESTs (Expression Sequence Tags) are publicly available, and the ESTs presumably cover not less than 80% of all human genes.

[0005] The information of ESTs is utilized for analyzing the structure of human genome, or for predicting the exonregions of genomic sequences or their expression profile. However, many human ESTs have been derived from proximal regions to the 3'-end of cDNA, and information around the 5'-end of mRNA is extremely little. Among these human
cDNAs, the number of the corresponding mRNAs whose encoding protein sequences are deduced is approximately
7000, and further, the number of full-length therein is only 5500. Thus, even including cDNA registered as EST, the
percentage of human cDNA obtained so far is estimated to be 10-15% of all the genes.

[0006] It is possible to identify the transcription start site of mRNA on the genomic sequence based on the 5'-end sequence of a full-length cDNA, and to analyze factors involved in the stability of mRNA that is contained in the cDNA, or in its regulation of expression at the translation stage. Also, since a full-length cDNA contains ATG, the translation start site, in the 5'-region, it can be translated into a protein in a correct frame. Therefore, it is possible to produce a large amount of the protein encoded by the cDNA or to analyze biological activity of the expressed protein by utilizing an appropriate expression system. Thus, analysis of a full-length cDNA provides valuable information which complements the information from genome sequencing. Also, full-length cDNA clones that can be expressed are extremely valuable in empirical analysis of gene function and in industrial application.

[0007] Therefore, if a novel human full-length cDNA is isolated, it can be used for developing medicines for diseases in which the gene is involved. The protein encoded by the gene can be used as a drug by itself. Thus, it has great significance to obtain a full-length cDNA encoding a novel human protein.

[0008] In particular, human secretory proteins or membrane proteins would be useful by itself as a medicine like tissue plasminogen activator (TPA), or as a target of medicines like membrane receptors. In addition, genes for signal transduction-associated proteins (protein kinases, etc.), glycoprotein-associated proteins, transcription-associated proteins, etc. are genes whose relationships to human diseases have been elucidated. Moreover, genes for disease-associated proteins form a gene group rich in genes whose relationships to human diseases have been elucidated.

[0009] Therefore, it has great significance to isolate novel full-length cDNA clones of human, only few of which has been isolated. Especially, isolation of a novel cDNA clone encoding a secretory protein or membrane protein is desired since the protein itself would be useful as a medicine, and also the clones potentially include a gene associated with diseases. In addition, genes encoding proteins that are associated with signal transduction, glycoprotein, transcription, or diseases are expected to be useful as target molecules for therapy, or as medicines themselves. These genes form a gene group predicted to be strongly associated with diseases. Thus, identification of the full-length cDNA clones

encoding those proteins has great significance.

#### SUMMARY OF THE INVENTION

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<sup>5</sup> [0010] An objective of the present invention is to provide a polynucleotide encoding a novel protein, a protein encoded by said polynucleotide, and novel usages of these.

[0011] The inventors have developed a method for efficiently cloning a human full-length cDNA that is predicted by the ATGpr etc. to be a full-length cDNA clone, from a full-length-enriched cDNA library that is synthesized by the oligocapping method. Then, the inventors determined the nucleotide sequence of the obtained cDNA clones from both 5'- and 3'- ends.

[0012] Furthermore, the inventors analyzed the obtained clones by the BLAST search of the databases, SwissProt (http://www.ebi.ac.uk/ebi\_docsSwissProt\_db/swisshome.html), GenBank (http://www.ncbi.nlm.nih.gov/web/GenBank), and UniGene (Human) (http://www.ncbi.nlm.nih.gov/UniGene).

[0013] The full-length cDNA clones of the present invention have high fullness ratio since these were obtained by the combination of (1) construction of a full-length-enriched cDNA library that is synthesized by the oligo-capping method, and (2) a system in which the full-length ratio is evaluated from the nucleotide sequence of the 5'-end (selection based on the ATGpr, previously removed complete sequences to ESTs). However, the primer of the present invention enables to obtain full-length cDNA easily without any specialized methods as in the described method.

Homology analysis in which the analysis is carried out against a not-full-length cDNA fragment to postulate the function of a protein encoded by said fragment, is being commonly performed.

However, since such analysis is based on the information of the fragment, it is not clear as to whether this fragment corresponds to a part that is functionally important in the protein. In other words, the reliability of the homology analysis based on the information of a fragment is doubtful, as information related to the structure of the whole protein is not available. However, the homology analysis of the present invention is conducted based on the information of a full-length cDNA comprising the whole coding region of the cDNA, and therefore, the homology of various portions of the protein can be analyzed. Hence, the reliability of the homology analysis has been dramatically improved in the present invention.

[0014] The inventors completed the invention by finding that it is possible to synthesize a novel full-length cDNA by using the combination of a primer that is designed based on the nucleotide sequence of the 5'-ends of the selected full-length cDNA clones and any of an oligo-dT primer or a 3'-primer that is designed based on the nucleotide sequence of the 3'-ends of the selected clones.

[0015] Thus, the present invention relates to primers described below, a method for synthesizing a polynucleotide using the primers, and polynucleotides obtained by the method.

[0016] First, the present invention relates to

(1) use of an oligonucleotide as a primer for synthesizing the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, or the complementary strand thereof, wherein said oligonucleotide is complementary to said polynucleotide or the complementary strand thereof and comprises at least 15 nucleotides;

(2) a primer set for synthesizing polynucleotides, the primer set comprising an oligo-dT primer and an oligonucleotide complementary to the complementary strand of the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, wherein said oligonucleotide comprises at least 15 nucleotides; and

(3) a primer set for synthesizing polynucleotides, the primer set comprising a combination of an oligonucleotide comprising a nucleotide sequence complementary to the complementary strand of the polynucleotide comprising a 5'-end nucleotide sequence and an oligonucleotide comprising a nucleotide sequence complementary to the polynucleotide comprising a 3'-end nucleotide sequence, wherein said oligonucleotides comprise at least 15 nucleotides and wherein said combination of 5'-end nucleotide sequence / 3'-end nucleotide sequence is selected from the combinations of 5'-end nucleotide sequence / 3'-end nucleotide sequence set forth in the SEQ ID NOs in Tables 1 and 2.

[0017] Tables 1 and 2 shows names of clones obtained in the examples described later, comprising the polynucleotide of the present invention (Table 1; 5547 clones, Table 2; 54 clones), names of nucleotide sequences at the 5'-end and 3'-end of the full-length cDNA, and their corresponding SEQ ID NOs. A blank indicates that the 3'-end sequence corresponding to the 5'-end sequence has not been determined for the same clone.

[0018] The SEQ ID NO of a 5'-end sequence is shown on the right side of the name of the 5'-end sequence, and the SEQ ID NO of a 3'-end sequence is shown on the right side of the name of the 3'-end sequence.

Table 1

nam	e of	name of	SEQ ID	name of	SEQ I	D
clo	ne	5'-end	of 5'-end	3'-end	of 3'-	-end
		sequence	sequence	sequence	seque	nce
HEM	BA1000005	F-HEMBA100000	5 1	R-HEMBA10000	05 5	548
HEM	BA1000012	F-HEMBA100001	2 2			
HEM	BA1000020	F-HEMBA100002	0 3			
HEM	BA1000030	F-HEMBA100003	0 4	R-HEMBA10000	30 5	549
HEM	BA1000042	F-HEMBA100004	2 5	R-HEMBA10000	42 5	550
HEM	BA1000046	F-HEMBA100004	6 6	R-HEMBA10000	46 5	551
HEM	BA1000050	F-HEMBA100005	0 7	R-HEMBA10000	50 5	552
HEM	BA1000076	F-HEMBA100007	6 8	R-HEMBA10000	76 5	553
HEM	BA1000111	F-HEMBA100011	1 9	R-HEMBA10001	.11 5	554
HEM	BA1000129	F-HEMBA100012	9 10	R-HEMBA10001	29 5	555
HEM	BA1000141	F-HEMBA100014	1 11	R-HEMBA10001	41 5	556
HEM	BA1000150	F-HEMBA100015	0 12	R-HEMBA10001	50 5	557
HEM	BA1000156	F-HEMBA100015	6 13	R-nnnnnnnnn	nn 5	558
HEM	BA1000158	F-HEMBA100015	8 14	R-HEMBA10001	.58 5	559
HEM	BA1000168	F-HEMBA100016	8 15	R-nnnnnnnn	nn 5	560

	HEMBA1000180	F-HEMBA1000180	16	R-HEMBA1000180	5561
	HEMBA1000185	F-HEMBA1000185	17	R-HEMBA1000185	5562
	HEMBA1000193	F-HEMBA1000193	18	R-HEMBA1000193	5563
5	HEMBA1000201	F-HEMBA1000201	19	R-HEMBA1000201	5564
	HEMBA1000213	F-HEMBA1000213	20	R-HEMBA1000213	5565
	HEMBA1000216	F-HEMBA1000216	21	R-HEMBA1000216	5566
	HEMBA1000227	F-HEMBA1000227	22	R-nnnnnnnnnnn	5567
10	HEMBA1000231	F-HEMBA1000231	23	R-HEMBA1000231	5568
	HEMBA1000243	F-HEMBA1000243	24	R-HEMBA1000243	5569
	HEMBA1000244	F-HEMBA1000244	25	R-HEMBA1000244	5570
	HEMBA1000251	F-HEMBA1000251	26	R-HEMBA1000251	5571
15	HEMBA1000264	F-HEMBA1000264	27	R-HEMBA1000264	5572
	HEMBA1000280	F-HEMBA1000280	28	R-nnnnnnnnnnn	5573
	HEMBA1000282	F-HEMBA1000282,	29	R-HEMBA1000282	5574
	HEMBA1000288	F-HEMBA1000288	30	R-HEMBA1000288	5575
20	HEMBA1000290	F-HEMBA1000290	31	R-HEMBA1000290	5576
	HEMBA1000302	F-HEMBA1000302	32	R-HEMBA1000302	5577
	HEMBA1000303	F-HEMBA1000303	33	R-nnnnnnnnnnn	5578
	HEMBA1000304	F-HEMBA1000304	34	R-nnnnnnnnnn	5579
25	HEMBA1000307	F-HEMBA1000307	35	R-HEMBA1000307	5580
25	HEMBA1000327	F-HEMBA1000327	36		
	HEMBA1000333	F-HEMBA1000333	37	R-nnnnnnnnnn	5581
	HEMBA1000338	F-HEMBA1000338	38	R-HEMBA1000338	5582
••	HEMBA1000351	F-HEMBA1000351	39	R-HEMBA1000351	5583
30	HEMBA1000355	F-HEMBA1000355	40	R-HEMBA1000355	5584
	HEMBA1000356	F-HEMBA1000356	41		
	HEMBA1000357	F-HEMBA1000357	42	R-HEMBA1000357	5585
	HEMBA1000366	F-HEMBA1000366	43	R-HEMBA1000366	5586
35	HEMBA1000369	F-HEMBA1000369	44	R-HEMBA1000369	5587
	HEMBA1000376	F-HEMBA1000376	45	R-HEMBA1000376	5588
	HEMBA1000387	F-HEMBA1000387	46	R-HEMBA1000387	5589
	HEMBA1000390	F-HEMBA1000390	47	R-HEMBA1000390	5590
40	HEMBA1000392	F-HEMBA1000392	48	R-HEMBA1000392	5591
	HEMBA1000396	F-HEMBA1000396	49	R-HEMBA1000396	5592
	HEMBA1000411	F-HEMBA1000411	50	R-HEMBA1000411	5593
	HEMBA1000418	F-HEMBA1000418	51	R-HEMBA1000418	5594
45	HEMBA1000422	F-HEMBA1000422	52	R-HEMBA1000422	5595
•	HEMBA1000428	F-HEMBA1000428	53	R-HEMBA1000428	5596
	HEMBA1000434	F-HEMBA1000434	54	R-HEMBA1000434	5597
	HEMBA1000442	F-HEMBA1000442	55	R-HEMBA1000442	5598
50	HEMBA1000456	F-HEMBA1000456	56	R-HEMBA1000456	5599
	HEMBA1000459	F-HEMBA1000459	57	R-HEMBA1000459	5600
	HEMBA1000460	F-HEMBA1000460	58	R-HEMBA1000460	5601
	HEMBA1000464	F-HEMBA1000464	59	R-HEMBA1000464	5602
55	HEMBA1000469	F-HEMBA1000469	60	R-HEMBA1000469	5603
	HEMBA1000488	F-HEMBA1000488	61	R-HEMBA1000488	5604

	HEMBA1000490	F-HEMBA1000490	62	R-HEMBA1000490	5605
	HEMBA1000491	F-HEMBA1000491	63	R-HEMBA1000491	5606
	HEMBA1000501	F-HEMBA1000501	64		
5	HEMBA1000504	F-HEMBA1000504	65	R-HEMBA1000504	5607
	HEMBA1000505	F-HEMBA1000505	66	R-HEMBA1000505	5608
	HEMBA1000508	F-HEMBA1000508	67	R-HEMBA1000508	5609
	HEMBA1000518	F-HEMBA1000518	68	R-HEMBA1000518	5610
10	HEMBA1000519	F-HEMBA1000519	69	R-HEMBA1000519	5611
	HEMBA1000520	F-HEMBA1000520	70	R-HEMBA1000520	5612
	HEMBA1000523	F-HEMBA1000523	71	R-HEMBA1000523	5613
	HEMBA1000531	F-HEMBA1000531	72	R-HEMBA1000531	5614
15	HEMBA1000534	F-HEMBA1000534	73	••	
.0	HEMBA1000540	F-HEMBA1000540	74	R-HEMBA1000540	5615
	HEMBA1000542	F-HEMBA1000542,	75	••••	
	HEMBA1000545	F-HEMBA1000545	76	R-HEMBA1000545	5616
20	HEMBA1000555	F-HEMBA1000555	77	R-nnnnnnnnnn	5617
20	HEMBA1000557	F-HEMBA1000557	78	R-HEMBA1000557	5618
	HEMBA1000561	F-HEMBA1000561	79	R-HEMBA1000561	5619
	HEMBA1000563	F-HEMBA1000563	80	R-HEMBA1000563	5620
	HEMBA1000568	F-HEMBA1000568	81	R-HEMBA1000568	5621
25	HEMBA1000569	F-HEMBA1000569	82	R-nnnnnnnnnnn	5622
	HEMBA1000575	F-HEMBA1000575	83	R-HEMBA1000575	5623
	HEMBA1000588	F-HEMBA1000588	84	R-HEMBA1000588	5624
	HEMBA1000591	F-HEMBA1000591	85	R-HEMBA1000591	5625
30	HEMBA1000592	F-HEMBA1000592	86	R-HEMBA1000592	5626
	HEMBA1000594	F-HEMBA1000594	87	R-HEMBA1000594	5627
	HEMBA1000604	F-HEMBA1000604	88	R-HEMBA1000604	5628
	HEMBA1000608	F-HEMBA1000608	89	R-HEMBA1000608	5629
35	HEMBA1000622	F-HEMBA1000622	90	R-HEMBA1000622	5630
	HEMBA1000636	F-HEMBA1000636	91	R-HEMBA1000636	5631
	HEMBA1000637	F-HEMBA1000637	92	R-HEMBA1000637	5632
	HEMBA1000655	F-HEMBA1000655	93	R-HEMBA1000655	5633
40	HEMBA1000657	F-HEMBA1000657	94	R-HEMBA1000657	5634
	HEMBA1000662	F-HEMBA1000662	95	R-HEMBA1000662	5635
	HEMBA1000673	F-HEMBA1000673	96	R-HEMBA1000673	5636
	HEMBA1000682	F-HEMBA1000682	97	R-HEMBA1000682	5637
45	HEMBA1000686	F-HEMBA1000686	98	R-HEMBA1000686	5638
	HEMBA1000702	F-HEMBA1000702	99	R-HEMBA1000702	5639
	HEMBA1000705	F-HEMBA1000705	100	R-HEMBA1000705	5640
	HEMBA1000719	F-HEMBA1000719	101	R-HEMBA1000719	5641
50	HEMBA1000722	F-HEMBA1000722	102	R-HEMBA1000722	5642
50	HEMBA1000726	F-HEMBA1000726	103	R-HEMBA1000726	5643
	HEMBA1000727	F-HEMBA1000727	104	R-HEMBA1000727	5644
	HEMBA1000747	F-HEMBA1000747	105	R-HEMBA1000747	5645
	HEMBA1000749		106	R-HEMBA1000749	5646
55	HEMBA1000752	F-HEMBA1000752	107	R-HEMBA1000752	5647

	HEMBA1000769	F-HEMBA1000769	108	R-HEMBA1000769	5648
	HEMBA1000773	F-HEMBA1000773	109	R-HEMBA1000773	5649
	HEMBA1000774	F-HEMBA1000774	110	R-HEMBA1000774	5650
5	HEMBA1000791	F-HEMBA1000791	111	R-HEMBA1000791	5651
	HEMBA1000817	F-HEMBA1000817	112	R-HEMBA1000817	5652
	HEMBA1000822	F-HEMBA1000822	113	R-HEMBA1000822	5653
	HEMBA1000827	F-HEMBA1000827	114	R-HEMBA1000827	5654
10	HEMBA1000843	F-HEMBA1000843	115	R-HEMBA1000843	5655
	HEMBA1000851	F-HEMBA1000851	116	R-HEMBA1000851	5656
	HEMBA1000852	F-HEMBA1000852	117	R-HEMBA1000852	5657
	HEMBA1000867	F-HEMBA1000867	118	R-HEMBA1000867	5658
15	HEMBA1000869	F-HEMBA1000869	119	R-HEMBA1000869	5659
	HEMBA1000870	F-HEMBA1000870	120	R-HEMBA1000870	5660
	HEMBA1000872	F-HEMBA1000872,	121	R-HEMBA1000872	5661
	HEMBA1000876	F-HEMBA1000876	122	R-HEMBA1000876	5662
20	HEMBA1000908	F-HEMBA1000908	123	R-HEMBA1000908	5663
20	HEMBA1000910	F-HEMBA1000910	124	R-HEMBA1000910	5664
	HEMBA1000918	F-HEMBA1000918	125	R-HEMBA1000918	5665
	HEMBA1000919	F-HEMBA1000919	126	R-HEMBA1000919	5666
	HEMBA1000934	F-HEMBA1000934	127	R-HEMBA1000934	5667
25	HEMBA1000942	F-HEMBA1000942	128	R-HEMBA1000942	5668
	HEMBA1000943	F-HEMBA1000943	129	R-HEMBA1000943	5669
	HEMBA1000946	F-HEMBA1000946	130	R-HEMBA1000946	5670
	HEMBA1000960	F-HEMBA1000960	131	R-HEMBA1000960	5671
30	HEMBA1000968	F-HEMBA1000968	132	R-HEMBA1000968	5672
	HEMBA1000971	F-HEMBA1000971	133	R-HEMBA1000971	5673
	HEMBA1000972	F-HEMBA1000972	134	R-HEMBA1000972	5674
	HEMBA1000974	F-HEMBA1000974	135	R-HEMBA1000974	5675
35	HEMBA1000975	F-HEMBA1000975	136	R-HEMBA1000975	5676
	HEMBA1000985	F-HEMBA1000985	137	R-HEMBA1000985	5677
	HEMBA1000986	F-HEMBA1000986	138	R-HEMBA1000986	5678
	HEMBA1000991	F-HEMBA1000991	139	R-HEMBA1000991	5679
40	HEMBA1001007	F-HEMBA1001007	140	R-HEMBA1001007	5680
	HEMBA1001008	F-HEMBA1001008	141	R-HEMBA1001008	5681
	HEMBA1001009	F-HEMBA1001009	142	R-HEMBA1001009	5682
	HEMBA1001017	F-HEMBA1001017	143	R-HEMBA1001017	5683
45	HEMBA1001019	F-HEMBA1001019	144	R-HEMBA1001019	5684
	HEMBA1001020	F-HEMBA1001020	145	R-HEMBA1001020	5685
	HEMBA1001022	F-HEMBA1001022	146	R-HEMBA1001022	5686
	HEMBA1001024	F-HEMBA1001024	147	R-HEMBA1001024	5687
50	HEMBA1001026	F-HEMBA1001026	148	R-HEMBA1001026	5688
30	HEMBA1001043	F-HEMBA1001043	149	R-nnnnnnnnnnn	5689
	HEMBA1001051	F-HEMBA1001051	150	R-HEMBA1001051	5690
	HEMBA1001052	F-HEMBA1001052	151	R-HEMBA1001052	5691
	HEMBA1001059	F-HEMBA1001059	152		
55	HEMBA1001060	F-HEMBA1001060	153	R-HEMBA1001060	5692

	·HEMBA1001071	F-HEMBA1001071	154	R-HEMBA1001071	5693
	HEMBA1001077	F-HEMBA1001077	155	R-HEMBA1001077	5694
	HEMBA1001080	F-HEMBA1001080	156	R-HEMBA1001080	5695
5	HEMBA1001085	F-HEMBA1001085	157	R-HEMBA1001085	5696
	HEMBA1001088	F-HEMBA1001088	158	R-HEMBA1001088	5697
	HEMBA1001094	F-HEMBA1001094	159	R-HEMBA1001094	5698
	HEMBA1001099	F-HEMBA1001099	160	R-HEMBA1001099	5699
10	HEMBA1001109	F-HEMBA1001109	161	R-HEMBA1001109	5700
	HEMBA1001121	F-HEMBA1001121	162	R-HEMBA1001121	5701
	HEMBA1001122	F-HEMBA1001122	163	R-HEMBA1001122	5702
	HEMBA1001123	F-HEMBA1001123	164	R-HEMBA1001123	5703
15	HEMBA1001133	F-HEMBA1001133	165	R-HEMBA1001133	5704
,,	HEMBA1001137	F-HEMBA1001137	166	R-HEMBA1001137	5705
	HEMBA1001140	F-HEMBA1001140,	167	R-HEMBA1001140	5706
	HEMBA1001172	F-HEMBA1001172	168	R-HEMBA1001172	5707
	HEMBA1001174	F-HEMBA1001174	169	R-HEMBA1001174	5708
20	HEMBA1001197	F-HEMBA1001197	170	R-HEMBA1001197	5709
	HEMBA1001208	F-HEMBA1001208	171	R-HEMBA1001208	5710
	HEMBA1001213	F-HEMBA1001213	172	•	
	HEMBA1001226	F-HEMBA1001226	173	R-HEMBA1001226	5711
25	HEMBA1001235	F-HEMBA1001235	174	R-HEMBA1001235	5712
	HEMBA1001247	F-HEMBA1001247	175	R-HEMBA1001247	5713
	HEMBA1001257	F-HEMBA1001257	176	R-HEMBA1001257	5714
	HEMBA1001265	F-HEMBA1001265	177	R-HEMBA1001265	5715
30	HEMBA1001281	F-HEMBA1001281	178	R-nnnnnnnnnnn	5716
	HEMBA1001286	F-HEMBA1001286	179	R-HEMBA1001286	5717
	HEMBA1001289	F-HEMBA1001289	180	R-HEMBA1001289	5718
	HEMBA1001294	F-HEMBA1001294	181	R-HEMBA1001294	5719
35	HEMBA1001299	F-HEMBA1001299	182	R-HEMBA1001299	5720
	HEMBA1001302	F-HEMBA1001302	183	R-HEMBA1001302	5721
	HEMBA1001303	F-HEMBA1001303	184	R-HEMBA1001303	5722
	HEMBA1001310	F-HEMBA1001310	185	R-HEMBA1001310	5723
40	HEMBA1001319	F-HEMBA1001319	186	R-HEMBA1001319	5724
	HEMBA1001323	F-HEMBA1001323	187	R-HEMBA1001323	5725
•	HEMBA1001326	F-HEMBA1001326	188	R-HEMBA1001326	5726
	HEMBA1001327	F-HEMBA1001327	189	R-HEMBA1001327	5727
45	HEMBA1001330	F-HEMBA1001330	190	R-HEMBA1001330	5728
	HEMBA1001351	F-HEMBA1001351	191	R-HEMBA1001351	5729
	HEMBA1001361	F-HEMBA1001361	192	R-HEMBA1001361	5730
	HEMBA1001375	F-HEMBA1001375	193	R-HEMBA1001375	5731
50	HEMBA1001377	F-HEMBA1001377	194	R-HEMBA1001377	5732
30	HEMBA1001383	F-HEMBA1001383	195	R-HEMBA1001383	5733
	HEMBA1001387	F-HEMBA1001387	196	R-HEMBA1001387	5734
	HEMBA1001388	F-HEMBA1001388	197	R-HEMBA1001388	5735
	HEMBA1001391	F-HEMBA1001391	198	R-HEMBA1001391	5736
55	HEMBA1001398	F-HEMBA1001398	199	R-HEMBA1001398	5737

	HEMBA1001405	F-HEMBA1001405	200	R-HEMBA1001405	5738
	HEMBA1001407	F-HEMBA1001407	201	R-HEMBA1001407	5739
	HEMBA1001411	F-HEMBA1001411	202	R-HEMBA1001411	5740
5	HEMBA1001413	F-HEMBA1001413	203	R-HEMBA1001413	5741
	HEMBA1001415	F-HEMBA1001415	204	R-HEMBA1001415	5742
	HEMBA1001432	F-HEMBA1001432	205	R-HEMBA1001432	5743
	HEMBA1001433	F-HEMBA1001433	206	R-HEMBA1001433	5744
10	HEMBA1001435	F-HEMBA1001435	207	R-HEMBA1001435	5745
	HEMBA1001442	F-HEMBA1001442	208	R-HEMBA1001442	5746
	HEMBA1001446	F-HEMBA1001446	209	R-HEMBA1001446	5747
	HEMBA1001450	F-HEMBA1001450	210	R-HEMBA1001450	5748
15	HEMBA1001454	F-HEMBA1001454	211	R-HEMBA1001454	5749
	HEMBA1001455	F-HEMBA1001455	212	R-HEMBA1001455	5750
	HEMBA1001463	F-HEMBA1001463,	213	R-HEMBA1001463	5751
	HEMBA1001476	F-HEMBA1001476	214	R-HEMBA1001476	5752
20	HEMBA1001478	F-HEMBA1001478	215	R-HEMBA1001478	5753
20	HEMBA1001497	F-HEMBA1001497	216	R-HEMBA1001497	5754
	HEMBA1001510	F-HEMBA1001510	217	R-HEMBA1001510	5755
	HEMBA1001515	F-HEMBA1001515	218	R-HEMBA1001515	5756
25	HEMBA1001517	F-HEMBA1001517	219	R-HEMBA1001517	5757
23	HEMBA1001522	F-HEMBA1001522	220	R-HEMBA1001522	5758
	HEMBA1001526	F-HEMBA1001526	221	R-HEMBA1001526	5759
	HEMBA1001533	F-HEMBA1001533	222	R-HEMBA1001533	5760
•	HEMBA1001557	F-HEMBA1001557	223	R-HEMBA1001557	5761
30	HEMBA1001566	F-HEMBA1001566	224	R-HEMBA1001566	5762
	HEMBA1001569	F-HEMBA1001569	225	R-HEMBA1001569	5763
	HEMBA1001570	F-HEMBA1001570	226	R-HEMBA1001570	5764
	HEMBA1001579	F-HEMBA1001579	227	R-HEMBA1001579	5765
35	HEMBA1001581	F-HEMBA1001581	228	R-HEMBA1001581	5766
	HEMBA1001585	F-HEMBA1001585	229	R-HEMBA1001585	5767 5760
	HEMBA1001589	F-HEMBA1001589	230	R-HEMBA1001589	5768 5760
	HEMBA1001595 HEMBA1001608	F-HEMBA1001595 F-HEMBA1001608	231 232	R-HEMBA1001595 R-HEMBA1001608	5769 5770
40	HEMBA1001620	F-HEMBA1001620	232	R-HEMBA1001620	5771
	HEMBA1001635	F-HEMBA1001635	234	R-nnnnnnnnnnn	5772
	HEMBA1001636	F-HEMBA1001636	235	R-HEMBA1001636	5773
	HEMBA1001640	F-HEMBA1001640	236	R-HEMBA1001640	5774
45	HEMBA1001647	F-HEMBA1001647	237	K ILLIIDIIIOO1040	0111
	HEMBA1001651	F-HEMBA1001651	238	R-nnnnnnnnnnn	5775
	HEMBA1001655	F-HEMBA1001655	239	R-HEMBA1001655	5776
	HEMBA1001658	F-HEMBA1001658	240	R-HEMBA1001658	5777
50	HEMBA1001661	F-HEMBA1001661	241	R-HEMBA1001661	5778
	HEMBA1001672	F-HEMBA1001672	242	R-HEMBA1001672	5779
	HEMBA1001675	F-HEMBA1001675	243	R-HEMBA1001675	5780
	HEMBA1001678	F-HEMBA1001678	244	R-HEMBA1001678	5781
55	HEMBA1001681	F-HEMBA1001681	245	R-HEMBA1001681	5782

	HEMBA1001702	F-HEMBA1001702	246	R-HEMBA1001702	5783
	HEMBA1001709	F-HEMBA1001709	247	R-HEMBA1001709	5784
	HEMBA1001711	F-HEMBA1001711	248	R-HEMBA1001711	5785
5	HEMBA1001712	F-HEMBA1001712	249	R-HEMBA1001712	5786
	HEMBA1001714	F-HEMBA1001714	250	R-HEMBA1001714	5787
	HEMBA1001718	F-HEMBA1001718	251	R-HEMBA1001718	5788
	HEMBA1001723	F-HEMBA1001723	252	R-HEMBA1001723	5789
10	HEMBA1001731	F-HEMBA1001731	253	R-HEMBA1001731	5790
	HEMBA1001734	F-HEMBA1001734	254	R-HEMBA1001734	5791
	HEMBA1001744	F-HEMBA1001744	255	R-HEMBA1001744	5792
	HEMBA1001745	F-HEMBA1001745	256	R-HEMBA1001745	5793
	HEMBA1001745	F-HEMBA1001746	257	R-HEMBA1001746	5794
15	HEMBA1001761	F-HEMBA1001761	258	R-HEMBA1001761	5795
	HEMBA1001781	F-HEMBA1001781,		R-HEMBA1001781	5796
	HEMBA1001784	F-HEMBA1001784	260	R-HEMBA1001784	5797
	HEMBA1001791	F-HEMBA1001791			5798
20			261	R-HEMBA1001791	
	HEMBA1001800 HEMBA1001803	F-HEMBA1001800 F-HEMBA1001803	262	R-HEMBA1001800	5799
			263	R-HEMBA1001803	5800
	HEMBA1001804 HEMBA1001808	F-HEMBA1001804 F-HEMBA1001808	264	R-nnnnnnnnnnn	5801
25	HEMBA1001809		265	R-HEMBA1001808	5802
		F-HEMBA1001809	266	R-HEMBA1001809	5803
	HEMBA1001815	F-HEMBA1001815	267	R-HEMBA1001815	5804
	HEMBA1001819	F-HEMBA1001819	268	R-HEMBA1001819	5805
30	HEMBA1001820	F-HEMBA1001820	269	R-HEMBA1001820	5806
55	HEMBA1001822	F-HEMBA1001822	270	R-nnnnnnnnnnnn	5807
	HEMBA1001824	F-HEMBA1001824	271	R-HEMBA1001824	5808
	HEMBA1001835	F-HEMBA1001835	272	R-HEMBA1001835	5809
	HEMBA1001844	F-HEMBA1001844	273	R-HEMBA1001844	5810
35	HEMBA1001847	F-HEMBA1001847	274	R-HEMBA1001847	5811
	HEMBA1001861	F-HEMBA1001861	275	R-HEMBA1001861	5812
	HEMBA1001864	F-HEMBA1001864	276	R-HEMBA1001864	5813
	HEMBA1001866	F-HEMBA1001866	277	R-HEMBA1001866	5814
40	HEMBA1001869	F-HEMBA1001869	278	R-nnnnnnnnnnn	5815
	HEMBA1001888	F-HEMBA1001888	279	R-HEMBA1001888	5816
	HEMBA1001896	F-HEMBA1001896	280	R-HEMBA1001896	5817
	HEMBA1001910	F-HEMBA1001910	281	R-HEMBA1001910	5818
45	HEMBA1001912	F-HEMBA1001912	282	R-HEMBA1001912	5819
	HEMBA1001913	F-HEMBA1001913	283	R-HEMBA1001913	5820
	HEMBA1001915	F-HEMBA1001915	284	R-HEMBA1001915	5821
	HEMBA1001918	F-HEMBA1001918	285	R-HEMBA1001918	5822
50	HEMBA1001921	F-HEMBA1001921	286	R-HEMBA1001921	5823
	HEMBA1001939	F-HEMBA1001939	287	R-HEMBA1001939	5824
	HEMBA1001940	F-HEMBA1001940	288	R-HEMBA1001940	5825
	HEMBA1001942	F-HEMBA1001942	289	R-HEMBA1001942	5826
	HEMBA1001945	F-HEMBA1001945	290	R-HEMBA1001945	5827
55	HEMBA1001950	F-HEMBA1001950	291	R-HEMBA1001950	5828

	HEMBA1001960	F-HEMBA1001960	292	R-HEMBA1001960	5829
	HEMBA1001962	F-HEMBA1001962	293	R-HEMBA1001962	5830
	HEMBA1001964	F-HEMBA1001964	294	R-HEMBA1001964	5831
5	HEMBA1001967	F-HEMBA1001967	295	R-HEMBA1001967	5832
	HEMBA1001979	F-HEMBA1001979	296	R-HEMBA1001979	5833
	HEMBA1001987	F-HEMBA1001987	297	R-HEMBA1001987	5834
	HEMBA1001991	F-HEMBA1001991	298	R-HEMBA1001991	5835
10	HEMBA1002003	F-HEMBA1002003	299	R-HEMBA1002003	5836
	HEMBA1002008	F-HEMBA1002008	300	R-HEMBA1002008	5837
	HEMBA1002018	F-HEMBA1002018	301	R-HEMBA1002018	5838
	HEMBA1002022	F-HEMBA1002022	302	R-HEMBA1002022	5839
	HEMBA1002035	F-HEMBA1002035	303	R-HEMBA1002035	5840
15	HEMBA1002039	F-HEMBA1002039	304	R-HEMBA1002039	5841
	HEMBA1002049	F-HEMBA1002049	305	R-HEMBA1002049	5842
	HEMBA1002084	F-HEMBA1002084	306	R-HEMBA1002084	5843
	HEMBA1002092	F-HEMBA1002092	307	R-HEMBA1002092	5844
20	HEMBA1002100	F-HEMBA1002100	308	R-HEMBA1002100	5845
	HEMBA1002102	F-HEMBA1002102	309	R-HEMBA1002102	5846
	HEMBA1002113	F-HEMBA1002113	310	R-HEMBA1002113	5847
	HEMBA1002119	F-HEMBA1002119	311	R-HEMBA1002119	5848
25	HEMBA1002125	F-HEMBA1002125	312	R-HEMBA1002125	5849
	HEMBA1002139	F-HEMBA1002139	313	R-HEMBA1002139	5850
	HEMBA1002144	F-HEMBA1002144	314	R-HEMBA1002144	5851
	HEMBA1002150	F-HEMBA1002150	315	R-HEMBA1002150	5852
30	HEMBA1002151	F-HEMBA 1002151	316	R-HEMBA1002151	5853
	HEMBA1002153	F-HEMBA1002153	317	R-HEMBA1002153	5854
	HEMBA1002160	F-HEMBA1002160	318	R-HEMBA1002160	5855
	HEMBA1002161	F-HEMBA1002161	319	R-HEMBA1002161	5856
35	HEMBA1002162	F-HEMBA1002162	320	R-HEMBA1002162	5857
30	HEMBA1002166	F-HEMBA1002166	321	R-HEMBA1002166	5858
	HEMBA1002177	F-HEMBA1002177	322	R-HEMBA1002177	5859
	HEMBA1002185	F-HEMBA1002185	323	R-HEMBA1002185	5860
	HEMBA1002189	F-HEMBA1002189	324	R-HEMBA1002189	5861
40	HEMBA1002191	F-HEMBA1002191	325	R-HEMBA1002191	5862
	HEMBA1002199	F-HEMBA1002199	326	R-HEMBA1002199	5863
	HEMBA1002204	F-HEMBA1002204	327	R-HEMBA1002204	5864
	HEMBA1002212	F-HEMBA1002212	328	R-HEMBA1002212	5865
45	HEMBA1002215	F-HEMBA1002215	329	R-HEMBA1002215	5866
	HEMBA1002226	F-HEMBA1002226	330	R-HEMBA1002226	5867
	HEMBA1002229	F-HEMBA1002229	331	R-HEMBA1002229	5868
	HEMBA1002237	F-HEMBA1002237	332	R-HEMBA1002237	5869
50	HEMBA1002241	F-HEMBA1002241	333		
	HEMBA1002253	F-HEMBA1002253	334	R-HEMBA1002253	5870
	HEMBA1002257	F-HEMBA1002257	335	R-HEMBA1002257	5871
	HEMBA1002265	F-HEMBA1002265	336		
55	HEMBA1002267	F-HEMBA1002267	337	R-HEMBA1002267	5872

	HEMBA1002270.	F-HEMBA1002270	338	R-HEMBA1002270	5873
	HEMBA1002321	F-HEMBA1002321	339	R-HEMBA1002321	5874
	HEMBA1002328		340	R-HEMBA1002328	5875
5	HEMBA1002337	F-HEMBA1002337	341	R-HEMBA1002337	5876
	HEMBA1002341	F-HEMBA1002341	342	R-HEMBA1002341	5877
	HEMBA1002348	F-HEMBA1002348	343	R-HEMBA1002348	5878
	HEMBA1002349	F-HEMBA1002349	344	R-HEMBA1002349	5879
10	HEMBA1002363	F-HEMBA1002363	345	R-nnnnnnnnnnn	5880
,,,	HEMBA1002381	F-HEMBA1002381	346	R-HEMBA1002381	5881
	HEMBA1002389	F-HEMBA1002389	347	R-HEMBA1002389	5882
	HEMBA1002417	F-HEMBA1002417	348	R-HEMBA1002417	5883
	HEMBA1002419	F-HEMBA1002419	349	R-HEMBA1002419	5884
15	HEMBA1002430	F-HEMBA1002430	350	R-HEMBA1002430	5885
	HEMBA1002439	F-HEMBA1002439,	351	R-HEMBA1002439	5886
	HEMBA1002458	F-HEMBA1002458	352	R-HEMBA1002458	5887
	HEMBA1002460	F-HEMBA1002460	353	R-HEMBA1002460	5888
20	HEMBA1002462	F-HEMBA1002462	354	R-HEMBA1002462	5889
	HEMBA1002469	F-HEMBA1002469	355		
	HEMBA1002475	F-HEMBA1002475	356	R-nnnnnnnnnnn	5890
	HEMBA1002477	F-HEMBA1002477	357	R-HEMBA1002477	5891
25	HEMBA1002486	F-HEMBA1002486	358	R-HEMBA1002486	5892
	HEMBA1002495	F-HEMBA1002495	359	R-HEMBA1002495	5893
	HEMBA1002498	F-HEMBA1002498	360	R-HEMBA1002498	5894
	HEMBA1002503	F-HEMBA1002503	361	R-HEMBA1002503	5895
30	HEMBA1002508	F-HEMBA1002508	362	R-HEMBA1002508	5896
	HEMBA1002513	F-HEMBA1002513	363	R-nnnnnnnnnnn	5897
	HEMBA1002515	F-HEMBA1002515	364	R-HEMBA1002515	5898
	HEMBA1002538	F-HEMBA1002538	365	R-HEMBA1002538	5899
35	HEMBA1002542	F-HEMBA1002542	366	R-HEMBA1002542	5900
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	HEMBA1002552	F-HEMBA1002552	368	R-HEMBA1002552	5902
	HEMBA1002555	F-HEMBA1002555	369	R-HEMBA1002555	5903
	HEMBA1002558	F-HEMBA1002558	370	R-HEMBA1002558	5904
40	HEMBA1002561	F-HEMBA1002561	371	R-HEMBA1002561	5905
	HEMBA1002569	F-HEMBA1002569	372	R-nnnnnnnnnnn	5906
	HEMBA1002583	F-HEMBA1002583	373	R-HEMBA1002583	5907
	HEMBA1002590	F-HEMBA1002590	374	R-HEMBA1002590	5908
45	HEMBA1002592	F-HEMBA1002592	375	R-HEMBA1002592	5909
	HEMBA1002609	F-HEMBA1002609	376		
	HEMBA1002621	F-HEMBA1002621	377	R-HEMBA1002621	5910
	HEMBA1002624	F-HEMBA1002624	378	R-HEMBA1002624	5911
50	HEMBA1002628	F-HEMBA1002628	379	R-HEMBA1002628	5912
	HEMBA1002629	F-HEMBA1002629	380	R-HEMBA1002629	5913
	HEMBA1002645	F-HEMBA1002645	381	R-HEMBA1002645	5914
	HEMBA1002651	F-HEMBA1002651	382	R-HEMBA1002651	5915
55	HEMBA1002659	F-HEMBA1002659	383	R-HEMBA1002659	5916

	HEMBA1002661	F-HEMBA1002661	384	R-HEMBA1002661	5917
	HEMBA1002666	F-HEMBA1002666	385	R-HEMBA1002666	5918
	HEMBA1002678	F-HEMBA1002678	386	R-HEMBA1002678	5919
5	HEMBA1002679	F-HEMBA1002679	387	R-nnnnnnnnnn	5920
	HEMBA1002688	F-HEMBA1002688	388	R-HEMBA1002688	5921
	HEMBA1002696	F-HEMBA1002696	389	R-HEMBA1002696	5922
	HEMBA1002703	F-HEMBA1002703	390		
10	HEMBA1002712	F-HEMBA1002712	391	R-HEMBA1002712	5923
	HEMBA1002716	F-HEMBA1002716	392	R-HEMBA1002716	5924
	HEMBA1002728	F-HEMBA1002728	393	R-HEMBA1002728	5925
	HEMBA1002730	F-HEMBA1002730	394	R-HEMBA1002730	5926
15	HEMBA1002742	F-HEMBA1002742	395	R-HEMBA1002742	5927
	HEMBA1002746	F-HEMBA1002746	396	R-HEMBA1002746	5928
	HEMBA1002748	F-HEMBA1002748,	397	R-HEMBA1002748	5929
	HEMBA1002750	F-HEMBA1002750	398	R-HEMBA1002750	5930
20	HEMBA1002768	F-HEMBA1002768	399	R-HEMBA1002768	5931
20	HEMBA1002770	F-HEMBA1002770	400	R-HEMBA1002770	5932
	HEMBA1002777	F-HEMBA1002777	401	R-HEMBA1002777	5933
	HEMBA1002779	F-HEMBA1002779	402	R-HEMBA1002779	5934
	HEMBA1002780	F-HEMBA1002780	403	R-HEMBA1002780	5935
25	HEMBA1002794	F-HEMBA1002794	404	R-HEMBA1002794	5936
	HEMBA1002801	F-HEMBA1002801	405	R-HEMBA1002801	5937
	HEMBA1002810	F-HEMBA1002810	406	R-HEMBA1002810	5938
	HEMBA1002816	F-HEMBA1002816	407	R-HEMBA1002816	5939
30	HEMBA1002818	F-HEMBA1002818	408		
	HEMBA1002826	F-HEMBA1002826	409	R-HEMBA1002826	5940
	HEMBA1002833	F-HEMBA1002833	410	R-HEMBA1002833	5941
	HEMBA1002850	F-HEMBA1002850	411	R-HEMBA1002850	5942
35	HEMBA1002863	F-HEMBA1002863	412	R-HEMBA1002863	5943
	HEMBA1002876	F-HEMBA1002876	413	R-HEMBA1002876	5944
	HEMBA1002886	F-HEMBA1002886	414	R-HEMBA1002886	5945
	HEMBA1002896	F-HEMBA1002896	415	R-HEMBA1002896	5946
40	HEMBA1002921	F-HEMBA1002921	416	R-HEMBA1002921	5947
-	HEMBA1002924	F-HEMBA1002924	417	R-HEMBA1002924	5948
	HEMBA1002934	F-HEMBA1002934	418	R-HEMBA1002934	5949
	HEMBA1002935	F-HEMBA1002935	419	R-HEMBA1002935	5950
	HEMBA1002937	F-HEMBA1002937	420	R-HEMBA1002937	5951
45	HEMBA1002939	F-HEMBA1002939	421	R-HEMBA1002939	5952
	HEMBA1002944	F-HEMBA1002944	422	R-HEMBA1002944	5953
	HEMBA1002951	F-HEMBA1002951	423	R-HEMBA1002951	5954
	HEMBA1002954	F-HEMBA1002954	424	R-HEMBA1002954	5955
50	HEMBA1002968	F-HEMBA1002968	425	R-HEMBA1002968	5956
	HEMBA1002970	F-HEMBA1002970	426	R-HEMBA1002970	5957
	HEMBA1002971	F-HEMBA1002971	427	R-HEMBA1002971	5958
	HEMBA1002973	F-HEMBA1002973	428	R-HEMBA1002973	5959
55	HEMBA1002997	F-HEMBA1002997	429	R-nnnnnnnnnnn	5960

	HEMBA1002999	F-HEMBA1002999	430	R-HEMBA1002999	5961
	HEMBA1003021	F-HEMBA1003021	431	R-HEMBA1003021	5962
	HEMBA1003033	F-HEMBA1003033	432	R-HEMBA1003033	5963
5	HEMBA1003034	F-HEMBA1003034	433	R-HEMBA1003034	5964
	HEMBA1003035	F-HEMBA1003035	434	R-HEMBA1003035	5965
	HEMBA1003037	F-HEMBA1003037	435	R-HEMBA1003037	5966
	HEMBA1003041	F-HEMBA1003041	436	R-HEMBA1003041	5967
10	HEMBA1003046	F-HEMBA1003046	437	R-HEMBA1003046	5968
	HEMBA1003064	F-HEMBA1003064	438	R-HEMBA1003064	5969
	HEMBA1003067	F-HEMBA1003067	439	R-HEMBA1003067	5970
	HEMBA1003071	F-HEMBA1003071	440	R-HEMBA1003071	5971
15	HEMBA1003077	F-HEMBA1003077	441	R-HEMBA1003077	5972
10	HEMBA1003078	F-HEMBA1003078	442	R-HEMBA1003078	5973
	HEMBA1003079	F-HEMBA1003079,	443	R-HEMBA1003079	5974
	HEMBA1003083	F-HEMBA1003083	444	R-HEMBA1003083	5975
	HEMBA1003086	F-HEMBA1003086	445	R-HEMBA1003086	5976
20	HEMBA1003096	F-HEMBA1003096	446	R-HEMBA1003096	5977
	HEMBA1003098	F-HEMBA1003098	447	R-HEMBA1003098	5978
	HEMBA1003117	F-HEMBA1003117	448	R-HEMBA1003117	5979
	HEMBA1003129	F-HEMBA1003129	449	R-HEMBA1003129	5980
25	HEMBA1003133	F-HEMBA1003133	450	R-HEMBA1003133	5981
	HEMBA1003136	F-HEMBA1003136	451	R-HEMBA1003136	5982
	HEMBA1003142	F-HEMBA1003142	452	R-HEMBA1003142	5983
	HEMBA1003148	F-HEMBA1003148	453	R-HEMBA1003148	5984
30	HEMBA1003166	F-HEMBA1003166	454	R-HEMBA1003166	5985
	HEMBA1003175	F-HEMBA1003175	455	R-HEMBA1003175	5986
	HEMBA1003179	F-HEMBA1003179	456		
	HEMBA1003197	F-HEMBA1003197	457	R-HEMBA1003197	5987
35	HEMBA1003199	F-HEMBA1003199	458	R-HEMBA1003199	5988
	HEMBA1003202	F-HEMBA1003202	459	R-HEMBA1003202	5989
	HEMBA1003204	F-HEMBA1003204	460	R-HEMBA1003204	5990
	HEMBA1003212	F-HEMBA1003212	461	R-HEMBA1003212	5991
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	HEMBA1003229	F-HEMBA1003229	464	R-HEMBA1003229	5994
	HEMBA1003235	F-HEMBA1003235	465	R-HEMBA1003235	5995
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45	HEMBA1003257	F-HEMBA1003257	467	R-HEMBA1003257	5997
	HEMBA1003273	F-HEMBA1003273	468	R-HEMBA1003273	5998
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50	HEMBA1003281	F-HEMBA1003281	471	R-HEMBA1003281	6001
	HEMBA1003286	F-HEMBA1003286	472		
	HEMBA1003291	F-HEMBA1003291	473	R-HEMBA1003291	6002
	HEMBA1003296	F-HEMBA1003296	474	R-HEMBA1003296	6003
55	HEMBA1003304	F-HEMBA1003304	475	R-HEMBA1003304	6004

	HEMBA1003309	F-HEMBA1003309	476	R-HEMBA1003309	6005
	HEMBA1003314	F-HEMBA1003314	477	R-HEMBA1003314	6006
	HEMBA1003322	F-HEMBA1003322	478	R-HEMBA1003322	6007
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	HEMBA1003328	F-HEMBA1003328	480	R-HEMBA1003328	6009
	HEMBA1003330	F-HEMBA1003330	481	R-HEMBA1003330	6010
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10	HEMBA1003369	F-HEMBA1003369	483	R-HEMBA1003369	6012
	HEMBA1003370	F-HEMBA1003370	484	R-HEMBA1003370	6013
	HEMBA1003373	F-HEMBA1003373	485	R-HEMBA1003373	6014
	HEMBA1003376	F-HEMBA1003376	486	R-HEMBA1003376	6015
15	HEMBA1003380	F-HEMBA1003380	487	R-HEMBA1003380	6016
10	HEMBA1003384	F-HEMBA1003384	488	R-HEMBA1003384	6017
	HEMBA1003395	F-HEMBA1003395,	489	R-HEMBA1003395	6018
	HEMBA1003402	F-HEMBA1003402	490	R-HEMBA1003402	6019
	HEMBA1003403	F-HEMBA1003403	491		
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	HEMBA1003417	F-HEMBA1003417	493	R-HEMBA1003417	6021
	HEMBA1003418	F-HEMBA1003418	494	R-HEMBA1003418	6022
	HEMBA1003433	F-HEMBA1003433	495	R-HEMBA1003433	6023
25	HEMBA1003447	F-HEMBA1003447	496		
	HEMBA1003461	F-HEMBA1003461	497	R-HEMBA1003461	6024
	HEMBA1003463	F-HEMBA1003463	498	R-HEMBA1003463	6025
	HEMBA1003480	F-HEMBA1003480	499	R-HEMBA1003480	6026
30	HEMBA1003528	F-HEMBA1003528	500	R-HEMBA1003528	6027
	HEMBA1003531	F-HEMBA1003531	501	R-HEMBA1003531	6028
	HEMBA1003538	F-HEMBA1003538	502	R-HEMBA1003538	6029
	HEMBA1003545	F-HEMBA1003545	503	R-HEMBA1003545	6030
35	HEMBA1003548	F-HEMBA1003548	504	R-HEMBA1003548	6031
33	HEMBA1003555	F-HEMBA1003555	505	R-HEMBA1003555	6032
	HEMBA1003556	F-HEMBA1003556	506	R-HEMBA1003556	6033
	HEMBA1003560	F-HEMBA1003560	507	R-HEMBA1003560	6034
	HEMBA1003568	F-HEMBA1003568	508	R-HEMBA1003568	6035
40	HEMBA1003569	F-HEMBA1003569	509	R-HEMBA1003569	6036
	HEMBA1003571	F-HEMBA1003571	510	R-HEMBA1003571	6037
	HEMBA1003579	F-HEMBA1003579	511	R-HEMBA1003579	6038
	HEMBA1003581	F-HEMBA1003581	512	R-HEMBA1003581	6039
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	HEMBA1003595	F-HEMBA1003595	514	R-HEMBA1003595	6041
	HEMBA1003597	F-HEMBA1003597	515	R-HEMBA1003597	6042
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	HEMBA1003621	F-HEMBA1003621	519	R-HEMBA1003621	6046
	HEMBA1003622	F-HEMBA1003622	520	R-HEMBA1003622	6047
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	HEMBA1003656	F-HEMBA1003656	526	R-HEMBA1003656	6053
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	HEMBA1003690	F-HEMBA1003690	532	R-HEMBA1003690	6059
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15	HEMBA1003711	F-HEMBA1003711	534	R-HEMBA1003711	6061
			535	R-HEMBA1003714	6062
	HEMBA1003714	F-HEMBA1003714,		R-HEMBA1003714	6063
	HEMBA1003715	F-HEMBA1003715	536		6064
20	HEMBA1003720	F-HEMBA1003720	537	R-HEMBA1003720	6065
	HEMBA1003725	F-HEMBA1003725	538	R-HEMBA1003725	
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25	HEMBA1003758	F-HEMBA1003758	542	R-HEMBA1003.758	6069
	HEMBA1003760	F-HEMBA1003760	543	R-HEMBA1003760	6070
	HEMBA1003773	F-HEMBA1003773	544	R-HEMBA1003773	6071
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	HEMBA1003804	F-HEMBA1003804	549	R-HEMBA1003804	6076
35	HEMBA1003805	F-HEMBA1003805	550	R-HEMBA1003805	6077
	HEMBA1003807	F-HEMBA1003807	551	R-HEMBA1003807	6078
	HEMBA1003827	F-HEMBA1003827	552		
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45	HEMBA1003880	F-HEMBA1003880	559	R-HEMBA1003880	6085
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	HEMBA1003939	F-HEMBA1003939	566	R-HEMBA1003939	6092
	HEMBA1003942	F-HEMBA1003942	567	R-HEMBA1003942	6093
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		HEMBA1003950	F-HEMBA1003950	568	R-HEMBA1003950	6094
		HEMBA1003953	F-HEMBA1003953	569	R-HEMBA1003953	6095
		HEMBA1003958	F-HEMBA1003958	570	R-HEMBA1003958	6096
;	5	HEMBA1003959	F-HEMBA1003959	571	R-HEMBA1003959	6097
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		HEMBA1003978	F-HEMBA1003978	573	R-HEMBA1003978	6099
		HEMBA1003985	F-HEMBA1003985	574	R-HEMBA1003985	6100
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		HEMBA1003989	F-HEMBA1003989	576	R-HEMBA1003989	6102
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		HEMBA1004024	F-HEMBA1004024	581	R-HEMBA1004024	6107
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		HEMBA1004042	F-HEMBA1004042	583	R-HEMBA1004042	6109
2	0	HEMBA1004045	F-HEMBA1004045	584	R-HEMBA1004045	6110
		HEMBA1004048	F-HEMBA1004048	585	R-HEMBA1004048	6111
		HEMBA1004049	F-HEMBA1004049	586	R-HEMBA1004049	6112
		HEMBA1004055	F-HEMBA1004055	587	R-HEMBA1004055	6113
2	25	HEMBA1004056	F-HEMBA1004056	588	R-HEMBA1004056	6114
		HEMBA1004074	F-HEMBA1004074	589	R-HEMBA1004074	6115
		HEMBA1004086	F-HEMBA1004086	590	R-HEMBA1004086	6116
		HEMBA1004097	F-HEMBA1004097	591	R-HEMBA1004097	6117
3	0	HEMBA1004111	F-HEMBA1004111	592		
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		HEMBA1004143	F-HEMBA1004143	597	R-HEMBA1004143	6122
		HEMBA1004146	F-HEMBA1004146	598	R-HEMBA1004146	6123
		HEMBA1004150	F-HEMBA1004150	599	R-HEMBA1004150	6124
4	0	HEMBA1004164 HEMBA1004168	F-HEMBA1004164	600	R-HEMBA1004164	6125
		HEMBA1004199	F-HEMBA1004168	601	R-HEMBA1004168	6126
		HEMBA1004199	F-HEMBA1004199 F-HEMBA1004200	602 603	R-HEMBA1004199	6127
		HEMBA1004200	F-HEMBA1004200	604	R-HEMBA1004200 R-HEMBA1004202	6128 6129
4	15	HEMBA1004202	F-HEMBA1004202	605	R-HEMBA1004202	6130
,	•	HEMBA1004207	F-HEMBA1004207	606	R-HEMBA1004207	6131
		HEMBA1004225	F-HEMBA1004225	607	R-HEMBA1004225	6132
		HEMBA1004227	F-HEMBA1004227	608	R-HEMBA1004227	6133
		HEMBA1004238	F-HEMBA1004238	609	R-HEMBA1004238	6134
5	50	HEMBA1004241	F-HEMBA1004241	610	R-HEMBA1004241	6135
		HEMBA1004246	F-HEMBA1004241	611	R-HEMBA1004241	6136
		HEMBA1004248	F-HEMBA1004248	612	R-HEMBA1004248	6137
		HEMBA1004264	F-HEMBA1004264	613	R-HEMBA1004264	6138
5	55		1 10000111004004	010	" ITHIDITION 201	0100

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	HEMBA1004274	F-HEMBA1004274	616		
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	HEMBA1004306	F-HEMBA1004306	622	R-HEMBA1004306	6146
	HEMBA1004312	F-HEMBA1004312	623	R-HEMBA1004312	6147
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	HEMBA1004327	F-HEMBA1004327	626	R-HEMBA1004327	6150
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30	HEMBA1004396	F-HEMBA1004405	639	R-HEMBA1004405	6163
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35	HEMBA1004460	F-HEMBA1004460	643	R-HEMBA1004460	6167
	HEMBA1004461	F-HEMBA1004461	644	R-HEMBA1004461	6168
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40	HEMBA1004499	F-HEMBA1004499	647	•	
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	HEMBA1004534	F-HEMBA1004534	652	R-HEMBA1004534	6175
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	HEMBA1004542	F-HEMBA1004542	654		
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55	HEMBA1004586	F-HEMBA1004586	659	R-HEMBA1004586	6181
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	HEMBA1004604	F-HEMBA1004604	661		
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5	HEMBA1004617	F-HEMBA1004617	663	R-HEMBA1004617	6184
	HEMBA1004629	F-HEMBA1004629	664	R-HEMBA1004629	6185
	HEMBA1004631	F-HEMBA1004631	665	R-HEMBA1004631	6186
	HEMBA1004632	F-HEMBA1004632	666	R-HEMBA1004632	6187
10	HEMBA1004637	F-HEMBA1004637	667	R-HEMBA1004637	6188
	HEMBA1004638	F-HEMBA1004638	668	R-HEMBA1004638	6189
	HEMBA1004666	F-HEMBA1004666	669	R-HEMBA1004666	6190
	HEMBA1004669	F-HEMBA1004669	670	R-HEMBA1004669	6191
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	HEMBA1004672	F-HEMBA1004672	672	R-HEMBA1004672	6193
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	HEMBA1004697	F-HEMBA1004697	674	R-HEMBA1004697	6195
20	HEMBA1004705	F-HEMBA1004705	675	R-HEMBA1004705	6196
	HEMBA1004709	F-HEMBA1004709	676	R-HEMBA1004709	6197
	HEMBA1004711	F-HEMBA1004711	677	R-HEMBA1004711 R-HEMBA1004725	6198 6199
	HEMBA1004725 HEMBA1004730	F-HEMBA1004725 F-HEMBA1004730	678 679	R-HEMBA1004730	6200
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20	HEMBA1004751	F-HEMBA1004751	684	R-HEMBA1004751	6205
30	HEMBA1004752	F-HEMBA1004752	685	R-HEMBA1004752	6206
	HEMBA1004753	F-HEMBA1004753	686	R-HEMBA1004753	6207
	HEMBA1004756	F-HEMBA1004756	687	R-HEMBA1004756	6208
	HEMBA1004758	F-HEMBA1004758	688	R-HEMBA1004758	6209
35	HEMBA1004763	F-HEMBA1004763	689	R-HEMBA1004763	6210
	HEMBA1004768	F-HEMBA1004768	690	R-HEMBA1004768	6211
	HEMBA1004770	F-HEMBA1004770	691	R-HEMBA1004770	6212
	HEMBA1004771	F-HEMBA1004771	692	R-HEMBA1004771	6213
40	HEMBA1004776	F-HEMBA1004776	693	R-HEMBA1004776	6214
	HEMBA1004778	F-HEMBA1004778	694	R-HEMBA1004778	6215
	HEMBA1004795	F-HEMBA1004795	695	R-nnnnnnnnnnn	6216
	HEMBA1004803	F-HEMBA1004803	696	R-HEMBA1004803	6217
45	HEMBA1004806	F-HEMBA1004806	697	R-HEMBA1004806 R-HEMBA1004807	6218 6219
	HEMBA1004807 HEMBA1004816	F-HEMBA1004807 F-HEMBA1004816	698 699	R-HEMBA1004816	6220
	HEMBA1004820	F-HEMBA1004820	700	R-HEMBA1004820	6221
	HEMBA1004847	F-HEMBA1004847	700	R-HEMBA1004847	6222
50	HEMBA1004850	F-HEMBA1004850	702	R-HEMBA1004850	6223
	HEMBA1004863	F-HEMBA1004863	703	R-HEMBA1004863	6224
	HEMBA1004864	F-HEMBA1004864	704	R-HEMBA1004864	6225
	HEMBA1004865	F-HEMBA1004865	705	R-HEMBA1004865	6226
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	HEMBA1004880	F-HEMBA1004880	706	R-HEMBA1004880	6227
	HEMBA1004889	F-HEMBA1004889	707	R-HEMBA1004889	6228
	HEMBA1004900	F-HEMBA1004900	708	R-HEMBA1004900	6229
5	HEMBA1004909	F-HEMBA1004909	709	R-HEMBA1004909	6230
	HEMBA1004918	F-HEMBA1004918	710	R-HEMBA1004918	6231
	HEMBA1004923	F-HEMBA1004923	711	R-HEMBA1004923	6232
	HEMBA1004929	F-HEMBA1004929	712	R-HEMBA1004929	6233
10	HEMBA1004930	F-HEMBA1004930	713	R-HEMBA1004930	6234
10	HEMBA1004933	F-HEMBA1004933	714	R-HEMBA1004933	6235
	HEMBA1004934	F-HEMBA1004934	715	R-HEMBA1004934	6236
	HEMBA1004944	F-HEMBA1004944	716	R-HEMBA1004944	6237
	HEMBA1004954	F-HEMBA1004954	717	R-HEMBA1004954	6238
15	HEMBA1004956	F-HEMBA1004956	718	R-HEMBA1004956	6239
	HEMBA1004960	F-HEMBA1004960,	719	R-HEMBA1004960	6240
	HEMBA1004972	F-HEMBA1004972	720	R-HEMBA1004972	6241
	HEMBA1004973	F-HEMBA1004973	721	R-HEMBA1004973	6242
20	HEMBA1004977	F-HEMBA1004977	722	R-HEMBA1004977	6243
	HEMBA1004978	F-HEMBA1004978	723	R-HEMBA1004978	6244
	HEMBA1004980	F-HEMBA1004980	724	R-HEMBA1004980	6245
	HEMBA1004983	F-HEMBA1004983	725	R-HEMBA1004983	6246
25	HEMBA1004995	F-HEMBA1004995	726	R-HEMBA1004995	6247
	HEMBA1005008	F-HEMBA1005008	727	R-HEMBA1005008	6248
	HEMBA1005009	F-HEMBA1005009	728	R-HEMBA1005009	6249
	HEMBA1005019	F-HEMBA1005019	729	R-HEMBA1005019	6250
30	HEMBA1005029	F-HEMBA1005029	730	R-HEMBA1005029	6251
30	HEMBA1005035	F-HEMBA1005035	731	R-HEMBA1005035	6252
	HEMBA1005039	F-HEMBA1005039	732	R-HEMBA1005039	6253
	HEMBA1005047	F-HEMBA1005047	733	R-HEMBA1005047	6254
	HEMBA1005050	F-HEMBA1005050	734	R-HEMBA1005050	6255
35	HEMBA1005062	F-HEMBA1005062	735	R-HEMBA1005062	6256
	HEMBA1005066	F-HEMBA1005066	736	R-HEMBA1005066	6257
	HEMBA1005075	F-HEMBA1005075	737	R-HEMBA1005075	6258
	HEMBA1005079	F-HEMBA1005079	738	R-HEMBA1005079	6259
40	HEMBA1005083	F-HEMBA1005083	739	R-HEMBA1005083	6260
	HEMBA1005101	F-HEMBA1005101	740	R-HEMBA1005101	6261
	HEMBA1005113	F-HEMBA1005113	741	R-HEMBA1005113	6262
	HEMBA1005123	F-HEMBA1005123	742	R-HEMBA1005123	6263
45	HEMBA1005133	F-HEMBA1005133	743	R-HEMBA1005133	6264
	HEMBA1005149	F-HEMBA1005149	744	R-HEMBA1005149	6265
	HEMBA1005152	F-HEMBA1005152	745	R-HEMBA1005152	6266
	HEMBA1005159	F-HEMBA1005159	746	R-HEMBA1005159	6267
50	HEMBA1005185	F-HEMBA1005185	747	R-HEMBA1005185	6268
50	HEMBA1005201	F-HEMBA1005201	748	. R-HEMBA1005201	6269
	HEMBA1005202	F-HEMBA1005202	749	R-HEMBA1005202	6270
	HEMBA1005206	F-HEMBA1005206	750		
	HEMBA1005219	F-HEMBA1005219	751	R-HEMBA1005219	6271

	HEMBA1005223	F-HEMBA1005223	752	R-HEMBA1005223	6272
	HEMBA1005232	F-HEMBA1005232	753	R-HEMBA1005232	6273
	HEMBA1005241	F-HEMBA1005241	754	R-HEMBA1005241	6274
5	HEMBA1005244	F-HEMBA1005244	755	R-HEMBA1005244	6275
	HEMBA1005251	F-HEMBA1005251	756	R-HEMBA1005251	6276
	HEMBA1005252	F-HEMBA1005252	757	R-HEMBA1005252	6277
	HEMBA1005274	F-HEMBA1005274	758	R-HEMBA1005274	6278
40	HEMBA1005275	F-HEMBA1005275	759	R-HEMBA1005275	6279
10	HEMBA1005293	F-HEMBA1005293	760	R-HEMBA1005293	6280
	HEMBA1005296	F-HEMBA1005296	761	R-HEMBA1005296	6281
	HEMBA1005304	F-HEMBA1005304	762	R-HEMBA1005304	6282
	HEMBA1005311	F-HEMBA1005311	763	R-HEMBA1005311	6283
15	HEMBA1005314	F-HEMBA1005314	764	R-HEMBA1005314	6284
	HEMBA1005315	F-HEMBA1005315,	765	R-HEMBA1005315	6285
	HEMBA1005318	F-HEMBA1005318	766	R-HEMBA1005318	6286
	HEMBA1005331	F-HEMBA1005331	767	R-HEMBA1005331	6287
20	HEMBA1005338	F-HEMBA1005338	768		
	HEMBA1005353	F-HEMBA1005353	769	R-HEMBA1005353	6288
	HEMBA1005359	F-HEMBA1005359	770	R-HEMBA1005359	6289
	HEMBA1005367	F-HEMBA1005367	771	R-HEMBA1005367	6290
25	HEMBA1005372	F-HEMBA1005372	772	R-HEMBA1005372	6291
	HEMBA1005374	F-HEMBA1005374	773	R-HEMBA1005374	6292
	HEMBA1005382	F-HEMBA1005382	774		
	HEMBA1005389	F-HEMBA1005389	775	R-HEMBA1005389	6293
30	HEMBA1005394	F-HEMBA1005394	776	R-HEMBA1005394	6294
30	HEMBA1005403	F-HEMBA1005403	777	R-HEMBA1005403	6295
	HEMBA1005408	F-HEMBA1005408	778	R-HEMBA1005408	6296
	HEMBA1005410	F-HEMBA1005410	779	R-HEMBA1005410	6297
	HEMBA1005411	F-HEMBA1005411	780	R-HEMBA1005411	6298
35	HEMBA1005423	F-HEMBA1005423	781	R-HEMBA1005423	6299
	HEMBA1005426	F-HEMBA1005426	782	R-HEMBA1005426	6300
	HEMBA1005443	F-HEMBA1005443	783	R-HEMBA1005443	6301
	HEMBA1005447	F-HEMBA1005447	784	R-HEMBA1005447	6302
40	HEMBA1005468	F-HEMBA1005468	785	R-HEMBA1005468	6303
	HEMBA1005469	F-HEMBA1005469	786	R-HEMBA1005469	6304
	HEMBA1005472	F-HEMBA1005472	787	R-HEMBA1005472	6305
	HEMBA1005474	F-HEMBA1005474	788		
45	HEMBA1005475	F-HEMBA1005475	789	R-HEMBA1005475	6306
	HEMBA1005497	F-HEMBA1005497	790	R-HEMBA1005497	6307
	HEMBA1005500	F-HEMBA1005500	791	R-HEMBA1005500	6308
	HEMBA1005506	F-HEMBA1005506	792	R-HEMBA1005506	6309
50	HEMBA1005508	F-HEMBA1005508	793	R-HEMBA1005508	6310
50	HEMBA1005511	F-HEMBA1005511	794	R-HEMBA1005511	6311
	HEMBA1005513	F-HEMBA1005513	795		
	HEMBA1005517	F-HEMBA1005517	796	R-HEMBA1005517	6312
	HEMBA1005518	F-HEMBA1005518	797	R-HEMBA1005518	6313

	HEMBA1005520	F-HEMBA1005520	798	R-HEMBA1005520	6314
	HEMBA1005526	F-HEMBA1005526	799	R-HEMBA1005526	6315
	HEMBA1005528	F-HEMBA1005528	800	R-HEMBA1005528	6316
5	HEMBA1005530	F-HEMBA1005530	801	R-HEMBA1005530	6317
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	HEMBA1005548	F-HEMBA1005548	802	R-HEMBA1005548	6318
	HEMBA1005552	F-HEMBA1005552	803	R-HEMBA1005552	6319
	HEMBA1005558	F-HEMBA1005558	804	R-HEMBA1005558	6320
10	HEMBA1005568	F-HEMBA1005568	805	R-HEMBA1005568	6321
	HEMBA1005570	F-HEMBA1005570	806	R-HEMBA1005570	6322
	HEMBA1005576	F-HEMBA1005576	807	R-HEMBA1005576	6323
	HEMBA1005577	F-HEMBA1005577	808	R-HEMBA1005577	6324
	HEMBA1005581	F-HEMBA1005581	809	R-HEMBA1005581	6325
15					6326
	HEMBA1005582	F-HEMBA1005582	810	R-HEMBA1005582	
	HEMBA1005583	F-HEMBA1005583,	811	R-HEMBA1005583	6327
	HEMBA1005588	F-HEMBA1005588	812	R-HEMBA1005588	6328
	HEMBA1005593	F-HEMBA1005593	813	R-HEMBA1005593	6329
20	HEMBA1005595	F-HEMBA1005595	814	R-HEMBA1005595	6330
	HEMBA1005606	F-HEMBA1005606	815	R-HEMBA1005606	6331
	HEMBA1005609	F-HEMBA1005609	816	R-HEMBA1005609	6332
	HEMBA1005616	F-HEMBA1005616	817	R-HEMBA1005616	6333
25	HEMBA1005621	F-HEMBA1005621	818	R-HEMBA1005621	6334
23	HEMBA1005627	F-HEMBA1005627	819	R-HEMBA1005627	6335
					6336
	HEMBA1005631	F-HEMBA1005631	820	R-HEMBA1005631	
	HEMBA1005632	F-HEMBA1005632	821	R-HEMBA1005632	6337
30	HEMBA1005634	F-HEMBA1005634	822	R-HEMBA1005634	6338
	HEMBA1005666	F-HEMBA1005666	823	R-HEMBA1005666	6339
	HEMBA1005670	F-HEMBA1005670	824	R-HEMBA1005670	6340
	HEMBA1005679	F-HEMBA1005679	825	R-HEMBA1005679	6341
	HEMBA1005680	F-HEMBA1005680	826	R-HEMBA1005680	6342
35	HEMBA1005685	F-HEMBA1005685	827	R-HEMBA1005685	6343
	HEMBA1005699	F-HEMBA1005699	828	R-HEMBA1005699	6344
	HEMBA1005705	F-HEMBA1005705	829	R-HEMBA1005705	6345
	HEMBA1005717	F-HEMBA1005717	830	R-HEMBA1005717	6346
40		F-HEMBA1005732	831	R-HEMBA1005732	6347
40	HEMBA1005732				
	HEMBA1005737	F-HEMBA1005737	832	R-HEMBA1005737	6348
	HEMBA1005746	F-HEMBA1005746	833	R-nnnnnnnnnnn	6349
	HEMBA1005755	F-HEMBA1005755	834	R-HEMBA1005755	6350
45	HEMBA1005765	F-HEMBA1005765	835	R-HEMBA1005765	6351
	HEMBA1005780	F-HEMBA1005780	836	R-HEMBA1005780	6352
	HEMBA1005813	F-HEMBA1005813	837	R-HEMBA1005813	6353
	HEMBA1005815	F-HEMBA1005815	838	R-HEMBA1005815	6354
	HEMBA1005822	F-HEMBA1005822	839	R-HEMBA1005822	6355
50	HEMBA1005829	F-HEMBA1005829	840	R-HEMBA1005829	6356
	HEMBA1005834	F-HEMBA1005834	841	R-HEMBA1005834	6357
		F-HEMBA1005852	842	R-HEMBA1005852	6358
	HEMBA1005852				
	HEMBA1005853	F-HEMBA1005853	843	R-HEMBA1005853	6359
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	VEIDA 100500A	E-UEVDA 1005994	011	D. HEMDATONEOOA	6260
	HEMBA1005884	F-HEMBA1005884	844	R-HEMBA1005884	6360
	HEMBA1005891	F-HEMBA1005891	845	R-HEMBA1005891	6361
	HEMBA1005894	F-HEMBA1005894	846	R-HEMBA1005894	6362
5	HEMBA1005909	F-HEMBA1005909	847	R-HEMBA1005909	6363
	HEMBA1005911	F-HEMBA1005911	848	R-HEMBA1005911	6364
	HEMBA1005921	F-HEMBA1005921	849	R-HEMBA1005921	6365
	HEMBA1005931	F-HEMBA1005931	850	R-HEMBA1005931	6366
10	HEMBA1005934	F-HEMBA1005934	851	R-HEMBA1005934	6367
	HEMBA1005962	F-HEMBA1005962	852	R-HEMBA1005962	6368
	HEMBA1005963	F-HEMBA1005963	853	R-HEMBA1005963	6369
	HEMBA1005990	F-HEMBA1005990	854	R-HEMBA1005990	6370
	HEMBA1005991	F-HEMBA1005991	855	R-HEMBA1005991	6371
15	HEMBA1005999	F-HEMBA1005999	856	R-HEMBA1005999	6372
	HEMBA1006002	F-HEMBA1006002,	857	R-HEMBA1006002	6373
	HEMBA1006005	F-HEMBA1006005	858	R-HEMBA1006005	6374
	HEMBA1006031	F-HEMBA1006031	859	R-nnnnnnnnnnn	6375
20	HEMBA1006035	F-HEMBA1006035	860	R-HEMBA1006035	6376
	HEMBA1006036	F-HEMBA1006036	861	R-HEMBA1006035	6377
				R-HEMBA1006042	6378
	HEMBA1006042	F-HEMBA1006042	862		6379
25	HEMBA1006067	F-HEMBA1006067	863	R-nnnnnnnnnnnn	6380
25	HEMBA1006081	F-HEMBA1006081	864	R-HEMBA1006081	
	HEMBA1006090	F-HEMBA1006090	865	R-HEMBA1006090	6381
	HEMBA1006091	F-HEMBA1006091	866	R-HEMBA1006091	6382
	HEMBA1006100	F-HEMBA1006100	867	R-HEMBA1006100	6383
30	HEMBA1006108	F-HEMBA1006108	868	R-HEMBA1006108	6384
	HEMBA1006121	F-HEMBA1006121	869	R-HEMBA1006121	6385
	HEMBA1006124	F-HEMBA1006124	870	R-HEMBA1006124	6386
	HEMBA1006130	F-HEMBA1006130	871	R-HEMBA1006130	6387
35	HEMBA1006138	F-HEMBA1006138	872	R-nnnnnnnnnnn	6388
33	HEMBA1006142	F-HEMBA1006142	873	R-HEMBA1006142	6389
	HEMBA1006155	F-HEMBA1006155	874	R-HEMBA1006155	6390
	HEMBA1006158	F-HEMBA1006158	875	R-HEMBA1006158	6391
	HEMBA1006173	F-HEMBA1006173	876	R-HEMBA1006173	6392
40	HEMBA1006182	F-HEMBA1006182	877	R-HEMBA1006182	6393
	HEMBA1006198	F-HEMBA1006198	878	R-HEMBA1006198	6394
	HEMBA1006235	F-HEMBA1006235	879	R-HEMBA1006235	6395
	HEMBA1006248	F-HEMBA1006248	880	R-HEMBA1006248	6396
45	HEMBA1006252	F-HEMBA1006252	881	R-HEMBA1006252	6397
	HEMBA1006253	F-HEMBA1006253	882	R-HEMBA1006253	6398
	HEMBA1006259	F-HEMBA1006259	883	R-HEMBA1006259	6399
	HEMBA1006268	F-HEMBA1006268	884	R-HEMBA1006268	6400
_	HEMBA1006272	F-HEMBA1006272	885	R-HEMBA1006272	6401
50	HEMBA1006278	F-HEMBA1006278	886	R-nnnnnnnnnn	6402
	HEMBA1006283	F-HEMBA1006283	887	R-HEMBA1006283	6403
	HEMBA1006284	F-HEMBA1006284	888	R-HEMBA1006284	6404
	HEMBA1006291	F-HEMBA1006291	889	R-HEMBA1006291	6405
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	HEMBA1006293	F-HEMBA1006293	890	R-HEMBA1006293	- 6406
	HEMBA1006309	F-HEMBA1006309	891	R-HEMBA1006309	6407
	HEMBA1006310	F-HEMBA1006310	892	R-HEMBA1006310	6408
5	HEMBA1006328	F-HEMBA1006328	893	R-HEMBA1006328	6409
	• • •	F-HEMBA1006334	894	R-HEMBA1006334	6410
	· ·	F-HEMBA1006344	895	R-HEMBA1006344	6411
	HEMBA1006347	F-HEMBA1006347	896	R-HEMBA1006347	6412
10	HEMBA1006349	F-HEMBA1006349	897	R-HEMBA1006349	6413
	HEMBA1006359	F-HEMBA1006359	898	R-HEMBA1006359	6414
	HEMBA1006364	F-HEMBA1006364	899	R-HEMBA1006364	6415
	HEMBA1006377	F-HEMBA1006377	900	R-HEMBA1006377	6416
	HEMBA1006380	F-HEMBA1006380	901	R-HEMBA1006380	6417
15	HEMBA1006381	F-HEMBA1006381	902	R-HEMBA1006381	6418
	HEMBA1006398	F-HEMBA1006398,	903	R-HEMBA1006398	6419
	HEMBA1006416	F-HEMBA1006416	904	R-HEMBA1006416	6420
	HEMBA1006419	F-HEMBA1006419	905	R-HEMBA1006419	6421
20	HEMBA1006421	F-HEMBA1006421	906	R-HEMBA1006421	6422
	HEMBA1006424	F-HEMBA1006424	907	R-HEMBA1006424	6423
	HEMBA1006426	F-HEMBA1006426	908	R-HEMBA1006426	6424
	HEMBA1006438	F-HEMBA1006438	909	R-HEMBA1006438	6425
25	HEMBA1006445	F-HEMBA1006445	910	R-HEMBA1006445	6426
	HEMBA1006446	F-HEMBA1006446	911	R-HEMBA1006446	6427
	HEMBA1006461	F-HEMBA1006461	912	R-HEMBA1006461	6428
	HEMBA1006467	F-HEMBA1006467	913	R-HEMBA1006467	6429
30	HEMBA1006471	F-HEMBA1006471	914	R-HEMBA1006471	6430
30	HEMBA1006474	F-HEMBA1006474	915	R-HEMBA1006474	6431
	HEMBA1006483	F-HEMBA1006483	916	R-HEMBA1006483	6432
	HEMBA1006485	F-HEMBA1006485	917	R-HEMBA1006485	6433
	HEMBA1006486	F-HEMBA1006486	918	R-HEMBA1006486	6434
35	HEMBA1006489	F-HEMBA1006489	919	R-HEMBA1006489	6435
	HEMBA1006492	F-HEMBA1006492	920	R-HEMBA1006492	6436
	HEMBA1006494	F-HEMBA1006494	921	R-HEMBA1006494	6437
	HEMBA1006497	F-HEMBA1006497	922	R-HEMBA1006497	6438
40	HEMBA1006502	F-HEMBA1006502	923	R-HEMBA1006502	6439
	HEMBA1006507	F-HEMBA1006507	924	R-HEMBA1006507	6440
	HEMBA1006521	F-HEMBA1006521	925	R-HEMBA1006521	6441
	HEMBA1006530	F-HEMBA1006530	926	R-HEMBA1006530	6442
45	HEMBA1006535	F-HEMBA1006535	927	R-HEMBA1006535	6443
	HEMBA1006540	F-HEMBA1006540	928	R-HEMBA1006540	6444
	HEMBA1006546	F-HEMBA1006546	929	R-HEMBA1006546	6445
	HEMBA1006559	F-HEMBA1006559	930	R-HEMBA1006559	6446
	HEMBA1006562	F-HEMBA1006562	931	R-HEMBA1006562	6447
50	HEMBA1006566	F-HEMBA1006566	932	R-HEMBA1006566	6448
	HEMBA1006569	F-HEMBA1006569	933	R-HEMBA1006569	6449
	HEMBA1006579	F-HEMBA1006579	934	R-HEMBA1006579	6450
	HEMBA1006583	F-HEMBA1006583	935	R-HEMBA1006583	6451

	HEMBA1006595	F-HEMBA1006595	936	R-HEMBA1006595	6452
	HEMBA1006597	F-HEMBA1006597	937	R-HEMBA1006597	6453
	HEMBA1006612	F-HEMBA1006612	938	R-HEMBA1006612	6454
5	HEMBA1006617	F-HEMBA1006617	939	R-nnnnnnnnnnn	6455
	HEMBA1006624	F-HEMBA1006624	940	R-HEMBA1006624	6456
	HEMBA1006631	F-HEMBA1006631	941	R-HEMBA1006631	6457
	HEMBA1006635	F-HEMBA1006635	942	R-HEMBA1006635	6458
10	HEMBA1006639	F-HEMBA1006639	943	R-HEMBA1006639	6459
10	HEMBA1006643	F-HEMBA1006643	944	R-HEMBA1006643	6460
	HEMBA1006648	F-HEMBA1006648	945	R-HEMBA1006648	6461
	HEMBA1006652	F-HEMBA1006652	946	R-HEMBA1006652	6462
	HEMBA1006653	F-HEMBA1006653	947	R-HEMBA1006653	6463
15	HEMBA1006659	F-HEMBA1006659	948		
	HEMBA1006665	F-HEMBA1006665,	949	R-HEMBA1006665	6464
	HEMBA1006674	F-HEMBA1006674	950	R-HEMBA1006674	6465
	HEMBA1006676	F-HEMBA1006676	951	R-HEMBA1006676	6466
20	HEMBA1006682	F-HEMBA1006682	952	R-HEMBA1006682	6467
	HEMBA1006695	F-HEMBA1006695	953	R-HEMBA1006695	6468
	HEMBA1006696	F-HEMBA1006696	954	R-HEMBA1006696	6469
	HEMBA1006708	F-HEMBA1006708	955	R-HEMBA1006708	6470
25	HEMBA1006709	F-HEMBA1006709	956	R-HEMBA1006709	6471
23	HEMBA1006717	F-HEMBA1006717	957	R-HEMBA1006717	6472
	HEMBA1006737	F-HEMBA1006737	958	R-HEMBA1006737	6473
	HEMBA1006744	F-HEMBA1006744	959	R-HEMBA1006744	6474
	HEMBA1006754	F-HEMBA1006754	960	R-HEMBA1006754	6475
30	HEMBA1006758	F-HEMBA1006758	961	R-HEMBA1006758	6476
	HEMBA1006767	F-HEMBA1006767	962	R-HEMBA1006767	6477
	HEMBA1006779	F-HEMBA1006779	963	R-HEMBA1006779	6478
	HEMBA1006780	F-HEMBA1006780	964	R-HEMBA1006780	6479
35	HEMBA1006789	F-HEMBA1006789	965	R-HEMBA1006789	6480
	HEMBA1006795	F-HEMBA1006795	966	R-HEMBA1006795	6481
	HEMBA1006796	F-HEMBA1006796	967	R-HEMBA1006796	6482
	HEMBA1006807	F-HEMBA1006807	968	R-HEMBA1006807	6483
40	HEMBA1006821	F-HEMBA1006821	969	R-HEMBA1006821	6484
	HEMBA1006824	F-HEMBA1006824	970	R-HEMBA1006824	6485
	HEMBA1006832	F-HEMBA1006832	971	R-HEMBA1006832	6486
	HEMBA1006849	F-HEMBA1006849	972	R-HEMBA1006849	6487
	HEMBA1006865	F-HEMBA1006865	973	R-HEMBA1006865	6488
45	HEMBA1006877	F-HEMBA1006877	974	R-nnnnnnnnnnn	6489
	HEMBA1006885	F-HEMBA1006885	975	R-HEMBA1006885	6490
	HEMBA1006900	F-HEMBA1006900	976	R-HEMBA1006900	6491
	HEMBA1006914	F-HEMBA1006914	977		
50	HEMBA1006921	F-HEMBA1006921	978	R-HEMBA1006921	6492
	HEMBA1006926	F-HEMBA1006926	979	R-HEMBA1006926	6493
	HEMBA1006929	F-HEMBA1006929	980	R-HEMBA1006929	6494
	HEMBA1006936	F-HEMBA1006936	981	R-HEMBA1006936	6495

	HEMBA1006938	F-HEMBA1006938	982	R-HEMBA1006938	6496
	HEMBA1006941	F-HEMBA1006941	983	R-HEMBA1006941	6497
	HEMBA1006949	F-HEMBA1006949	984	R-HEMBA1006949	6498
5	HEMBA1006973	F-HEMBA1006973	985	R-HEMBA1006973	6499
	HEMBA1006976	F-HEMBA1006976	986	R-HEMBA1006976	6500
	HEMBA1006993		987	R-HEMBA1006993	6501
	HEMBA1006996		988	R-HEMBA1006996	6502
10	HEMBA1007002		989	R-HEMBA1007002	6503
	HEMBA1007017		990	R-HEMBA1007017	6504
	HEMBA1007018		991	R-HEMBA1007018	6505
	HEMBA1007045		992	R-HEMBA1007045	6506
	HEMBA1007051		993	R-HEMBA1007051	6507
15	HEMBA1007052		994	R-HEMBA1007052	6508
	HEMBA1007062			R-HEMBA1007062	6509
	HEMBA1007066		996	R-HEMBA1007066	6510
	HEMBA1007073		997	R-HEMBA1007073	6511
20	HEMBA1007078		998	R-HEMBA1007078	6512
	HEMBA1007080	•	999		
	HEMBA1007085	F-HEMBA1007085	1000	R-HEMBA1007085	6513
	HEMBA1007087	F-HEMBA1007087	1001	R-HEMBA1007087	6514
25	HEMBA1007112	F-HEMBA1007112	1002	R-HEMBA1007112	6515
	HEMBA1007113	F-HEMBA1007113	1003	R-HEMBA1007113	6516
	HEMBA1007121	F-HEMBA1007121	1004		
	HEMBA1007129	F-HEMBA1007129	1005	R-HEMBA1007129	6517
•	HEMBA1007147	7 F-HEMBA1007147	1006	R-HEMBA1007147	6518
30	HEMBA1007149	F-HEMBA1007149	1007	R-HEMBA1007149	6519
	HEMBA1007151	F-HEMBA1007151	1008	R-HEMBA1007151	6520
	HEMBA1007174	4 F-HEMBA1007174	1009	R-nnnnnnnnnnn	6521
	HEMBA1007178	F-HEMBA1007178	1010	R-HEMBA1007178	6522
35	HEMBA1007194	4 F-HEMBA1007194	1011	R-HEMBA1007194	6523
	HEMBA1007203	B F-HEMBA1007203	1012	R-HEMBA1007203	6524
	HEMBA1007206	6 F-HEMBA1007206	1013	R-HEMBA1007206	6525
	HEMBA1007224	4 F-HEMBA1007224	1014	R-HEMBA1007224	6526
40	HEMBA1007243	3 F-HEMBA1007243	1015		
	HEMBA100725		1016	R-HEMBA1007251	6527
	HEMBA100725	6 F-HEMBA1007256	1017	R-HEMBA1007256	6528
	HEMBA100726	7 F-HEMBA1007267	1018	R-HEMBA1007267	6529
45	HEMBA100727		1019	R-HEMBA1007273	6530
73	HEMBA1007279	9 F-HEMBA1007279	1020	R-HEMBA1007279	6531
	HEMBA100728		1021	R-HEMBA1007281	6532
	HEMBA100728		1022	R-HEMBA1007288	6533
	HEMBA100730		1023	R-HEMBA1007300	6534
50	HEMBA100730		1024	R-HEMBA1007301	6535
	HEMBA100731		1025	R-HEMBA1007319	6536
	HEMBA100732		1026	R-HEMBA1007320	6537
	HEMBA100732	2 F-HEMBA1007322	1027	R-HEMBA1007322	6538

	HEMBA1007327	F-HEMBA1007327	1028	R-HEMBA1007327	6539
	HEMBA1007341	F-HEMBA1007341	1029	R-HEMBA1007341	6540
	HEMBA1007342	F-HEMBA1007342	1030	R-HEMBA1007342	6541
5	HEMBA1007347	F-HEMBA1007347	1031	R-HEMBA1007347	6542
	HEMBB1000005	F-HEMBB1000005	1032	R-HEMBB1000005	6543
	HEMBB1000008	F-HEMBB1000008	1033	R-HEMBB1000008	6544
	HEMBB1000018	F-HEMBB1000018	1034	R-HEMBB1000018	6545
10	HEMBB1000024	F-HEMBB1000024	1035	R-HEMBB1000024	6546
10	HEMBB1000025	F-HEMBB1000025	1036	R-HEMBB1000025	6547
	HEMBB1000030	F-HEMBB1000030	1037	R-HEMBB1000030	6548
	HEMBB1000036	F-HEMBB1000036	1038	R-HEMBB1000036	6549
	HEMBB1000037	F-HEMBB1000037	1039	R-HEMBB1000037	6550
15	HEMBB1000039	F-HEMBB1000039	1040	R-HEMBB1000039	6551
	HEMBB1000044	F-HEMBB1000044,	1041	R-HEMBB1000044	6552
	HEMBB1000048	F-HEMBB1000048	1042	R-HEMBB1000048	6553
	HEMBB1000050	F-HEMBB1000050	1043	R-HEMBB1000050	6554
20	HEMBB1000054	F-HEMBB1000054	1044	R-HEMBB1000054	6555
	HEMBB1000055	F-HEMBB1000055	1045	R-HEMBB1000055	6556
	HEMBB1000059	F-HEMBB1000059	1046	R-HEMBB1000059	6557
	HEMBB1000083	F-HEMBB1000083	1047	R-HEMBB1000083	6558
25	HEMBB1000089	F-HEMBB1000089	1048	R-HEMBB1000089	6559
	HEMBB1000099	F-HEMBB1000099	1049	R-HEMBB1000099	6560
	HEMBB1000103	F-HEMBB1000103	1050	R-HEMBB1000103	6561
	HEMBB1000113	F-HEMBB1000113	1051	R-HEMBB1000113	6562
	HEMBB1000119	F-HEMBB1000119	1052	R-HEMBB1000119	6563
30	HEMBB1000136	F-HEMBB1000136	1053	R-HEMBB1000136	6564
	HEMBB1000141	F-HEMBB1000141	1054	R-HEMBB1000141	6565
	HEMBB1000144	F-HEMBB1000144	1055	R-HEMBB1000144	6566
	HEMBB1000173	F-HEMBB1000173	1056	R-HEMBB1000173	6567
35	HEMBB1000175	F-HEMBB1000175	1057	R-HEMBB1000175	6568
	HEMBB1000198	F-HEMBB1000198	1058	R-HEMBB1000198	6569
	HEMBB1000215	F-HEMBB1000215	1059	R-HEMBB1000215	6570
	HEMBB1000217	F-HEMBB1000217	1060	R-HEMBB1000217	6571
40	HEMBB1000218	F-HEMBB1000218	1061	R-HEMBB1000218	6572
	HEMBB1000226	F-HEMBB1000226	1062	R-HEMBB1000226	6573
	HEMBB1000240	F-HEMBB1000240	1063	R-HEMBB1000240	6574
	HEMBB1000244	F-HEMBB1000244	1064	R-HEMBB1000244	6575
45	HEMBB1000250	F-HEMBB1000250	1065	R-HEMBB1000250	6576
40	HEMBB1000258	F-HEMBB1000258	1066	R-HEMBB1000258	6577
	HEMBB1000264	F-HEMBB1000264	1067	R-HEMBB1000264	6578
	HEMBB1000266	F-HEMBB1000266	1068	R-HEMBB1000266	6579
	HEMBB1000272	F-HEMBB1000272	1069	R-HEMBB1000272	6580
50	HEMBB1000274	F-HEMBB1000274	1070	R-HEMBB1000274	6581
	HEMBB1000284	F-HEMBB1000284	1071	R-HEMBB1000284	6582
	HEMBB1000307	F-HEMBB1000307	1072	R-HEMBB1000307	6583
	HEMBB1000312	F-HEMBB1000312	1073	R-HEMBB1000312	6584

	HEMBB1000317	F-HEMBB1000317	1074	R-HEMBB1000317	6585
	HEMBB1000318	F-HEMBB1000318	1075	R-HEMBB1000318	6586
	HEMBB1000335	F-HEMBB1000335	1076	R-HEMBB1000335	6587
5	HEMBB1000336	F-HEMBB1000336	1077	R-HEMBB1000336	6588
	HEMBB1000337	F-HEMBB1000337	1078	R-HEMBB1000337	6589
	HEMBB1000338	F-HEMBB1000338	1079	R-HEMBB1000338	6590
	HEMBB1000339	F-HEMBB1000339	1080	R-HEMBB1000339	6591
10	HEMBB1000341	F-HEMBB1000341	1081	R-HEMBB1000341	6592
	HEMBB1000343	F-HEMBB1000343	1082	R-HEMBB1000343	6593
	HEMBB1000354	F-HEMBB1000354	1083	R-HEMBB1000354	6594
	HEMBB1000369	F-HEMBB1000369	1084	R-HEMBB1000369	6595
15	HEMBB1000374	F-HEMBB1000374	1085	R-HEMBB1000374	6596
13	HEMBB1000376	F-HEMBB1000376	1086	R-HEMBB1000376	6597
	HEMBB1000391	F-HEMBB1000391,	1087	R-HEMBB1000391	6598
	HEMBB1000399	F-HEMBB1000399	1088	R-HEMBB1000399	6599
	HEMBB1000402	F-HEMBB1000402	1089	R-HEMBB1000402	6600
20	HEMBB1000404	F-HEMBB1000404	1090	R-HEMBB1000404	6601
	HEMBB1000420	F-HEMBB1000420	1091	R-HEMBB1000420	6602
	HEMBB1000434	F-HEMBB1000434	1092	R-HEMBB1000434	6603
	HEMBB1000438	F-HEMBB1000438	1093	R-HEMBB1000438	6604
25	HEMBB1000441	F-HEMBB1000441	1094	R-HEMBB1000441	6605
	HEMBB1000449	F-HEMBB1000449	1095	R-HEMBB1000449	6606
	HEMBB1000455	F-HEMBB1000455	1096	R-HEMBB1000455	6607
	HEMBB1000472	F-HEMBB1000472	1097	R-HEMBB1000472	6608
30	HEMBB1000480	F-HEMBB1000480	1098	R-HEMBB1000480	6609
	HEMBB1000487	F-HEMBB1000487	1099	R-HEMBB1000487	6610
	HEMBB1000490	F-HEMBB1000490	1100	R-HEMBB1000490	6611
	HEMBB1000491	F-HEMBB1000491	1101	R-HEMBB1000491	6612
	HEMBB1000493	F-HEMBB1000493	1102	R-HEMBB1000493	6613
35	HEMBB1000510	F-HEMBB1000510	1103	R-HEMBB1000510	6614
	HEMBB1000518	F-HEMBB1000518	1104	R-HEMBB1000518	6615
	HEMBB1000523	F-HEMBB1000523	1105	R-HEMBB1000523	6616
	HEMBB1000530	F-HEMBB1000530	1106	R-HEMBB1000530	6617
40	HEMBB1000550	F-HEMBB1000550	1107	R-HEMBB1000550	6618
	HEMBB1000554	F-HEMBB1000554	1108	R-HEMBB1000554	6619
	HEMBB1000556	F-HEMBB1000556	1109	R-HEMBB1000556	6620
	HEMBB1000564	F-HEMBB1000564	1110	R-HEMBB1000564	6621
45	HEMBB1000573	F-HEMBB1000573	1111	R-HEMBB1000573	6622
	HEMBB1000575	F-HEMBB1000575	1112	R-HEMBB1000575	6623
	HEMBB1000586	F-HEMBB1000586	1113	R-HEMBB1000586	6624
	HEMBB1000589	F-HEMBB1000589	1114	R-HEMBB1000589	6625
50	HEMBB1000591	F-HEMBB1000591	1115	R-HEMBB1000591	6626
50	HEMBB1000592	F-HEMBB1000592	1116	R-HEMBB1000592	6627
	HEMBB1000593	F-HEMBB1000593	1117		
	HEMBB1000598	F-HEMBB1000598	1118	R-HEMBB1000598	6628
	HEMBB1000623	F-HEMBB1000623	1119	R-HEMBB1000623	6629

		HEMBB1000630	F-HEMBB1000630	1120	R-HEMBB1000630	6630
		HEMBB1000631	F-HEMBB1000631	1121	R-HEMBB1000631	6631
		HEMBB1000632	F-HEMBB1000632	1122	R-HEMBB1000632	6632
	5	HEMBB1000637	F-HEMBB1000637	1123	R-HEMBB1000637	6633
		HEMBB1000638	F-HEMBB1000638	1124	R-HEMBB1000638	6634
		HEMBB1000643	F-HEMBB1000643	1125	R-HEMBB1000643	6635
		HEMBB1000649	F-HEMBB1000649	1126	R-HEMBB1000649	6636
1	0	HEMBB1000652	F-HEMBB1000652	1127	R-HEMBB1000652	6637
		HEMBB1000665	F-HEMBB1000665	1128	R-HEMBB1000665	6638
		HEMBB1000671	F-HEMBB1000671	1129	R-HEMBB1000671	6639
		HEMBB1000673	F-HEMBB1000673	1130	R-HEMBB1000673	6640
1	5	HEMBB1000684	F-HEMBB1000684	1131	R-HEMBB1000684	6641
		HEMBB1000693	F-HEMBB1000693	1132	R-nnnnnnnnnnn	6642
		HEMBB1000705	F-HEMBB1000705	1133	R-HEMBB1000705	6643
		HEMBB1000706	F-HEMBB1000706	1134	R-HEMBB1000706	6644
,	ro	HEMBB1000709	F-HEMBB1000709	1135	R-HEMBB1000709	6645
-	<b>U</b>	HEMBB1000725	F-HEMBB1000725	1136	R-HEMBB1000725	6646
		HEMBB1000726	F-HEMBB1000726	1137	R-HEMBB1000726	6647
		HEMBB1000738	F-HEMBB1000738	1138	R-HEMBB1000738	6648
		HEMBB1000749	F-HEMBB1000749	1139	R-HEMBB1000749	6649
2	5	HEMBB1000763	F-HEMBB1000763	1140	R-HEMBB1000763	6650
		HEMBB1000770	F-HEMBB1000770	1141	R-HEMBB1000770	6651
		HEMBB1000774	F-HEMBB1000774	1142		
		HEMBB1000781	F-HEMBB1000781	1143	R-HEMBB1000781	6652
3	0	HEMBB1000789	F-HEMBB1000789	1144	R-HEMBB1000789	6653
		HEMBB1000790	F-HEMBB1000790	1145	R-HEMBB1000790	6654
		HEMBB1000794	F-HEMBB1000794	1146	R-HEMBB1000794	6655
		HEMBB1000807	F-HEMBB1000807	1147	R-HEMBB1000807	6656
3	5	HEMBB1000810	F-HEMBB1000810	1148	R-HEMBB1000810	6657
		HEMBB1000821	F-HEMBB1000821	1149	R-HEMBB1000821	6658
		HEMBB1000822	F-HEMBB1000822	1150	R-HEMBB1000822	6659
		HEMBB1000826	F-HEMBB1000826	1151	R-HEMBB1000826	6660
4	o	HEMBB1000827	F-HEMBB1000827	1152	R-HEMBB1000827	6661
		HEMBB1000831	F-HEMBB1000831	1153	R-HEMBB1000831	6662
		HEMBB1000835	F-HEMBB1000835	1154	R-HEMBB1000835	6663
		HEMBB1000840	F-HEMBB1000840	1155	R-HEMBB1000840	6664
4	5	HEMBB1000848	F-HEMBB1000848	1156	R-HEMBB1000848	6665
		HEMBB1000852	F-HEMBB1000852	1157	R-HEMBB1000852	6666
		HEMBB1000870	F-HEMBB1000870	1158	R-HEMBB1000870	6667
		HEMBB1000876	F-HEMBB1000876	1159	R-HEMBB1000876	6668
5	0	HEMBB1000883	F-HEMBB1000883	1160	R-HEMBB1000883	6669
J	•	HEMBB1000887	F-HEMBB1000887	1161	R-HEMBB1000887	6670
		HEMBB1000888	F-HEMBB1000888	1162	R-HEMBB1000888	6671
		HEMBB1000890	F-HEMBB1000890	1163	R-HEMBB1000890	6672
	_	HEMBB1000893	F-HEMBB1000893	1164	R-HEMBB1000893	6673
5.	•	HEMBB1000908	F-HEMBB1000908	1165	R-HEMBB1000908	6674

	HEMBB100091	0 F-HEMBB1000910	1166	R-HEMBB1000910	6675
	HEMBB100091	3 F-HEMBB1000913	1167	R-HEMBB1000913	6676
	HEMBB100091	5 F-HEMBB1000915	1168	R-HEMBB1000915	6677
5	HEMBB100091	7 F-HEMBB1000917	1169	R-HEMBB1000917	6678
	HEMBB100092	•		R-HEMBB1000927	6679
	HEMBB100094			R-HEMBB1000947	6680
	HEMBB100095			R-HEMBB1000959	6681
10	HEMBB100097			R-HEMBB1000973	6682
	HEMBB100097			R-HEMBB1000975	6683
	HEMBB100098			R-HEMBB1000981	6684
	HEMBB100098	5 F-HEMBB1000985	1176	R-HEMBB1000985	6685
15	HEMBB100099			R-HEMBB1000991	6686
	HEMBB100099			R-HEMBB1000996	6687
	HEMBB100100			R-HEMBB1001004	6688
	HEMBB100100		,	R-HEMBB1001008	6689
20	HEMBB100101	1 F-HEMBB1001011		R-HEMBB1001011	6690
20	HEMBB100101			R-HEMBB1001014	6691
	HEMBB100102	0 F-HEMBB1001020	1183	R-HEMBB1001020	6692
	HEMBB100102	4 F-HEMBB1001024	1184	R-HEMBB1001024	6693
05	HEMBB100103	7 F-HEMBB1001037	1185	R-HEMBB1001037	6694
25	HEMBB100104	7 F-HEMBB1001047	1186	R-HEMBB1001047	6695
	HEMBB100105	1 F-HEMBB1001051	1187	R-HEMBB1001051	6696
	HEMBB100105	6 F-HEMBB1001056	1188	R-HEMBB1001056	6697
	HEMBB100105	8 F-HEMBB1001058	1189	R-HEMBB1001058	6698
30	HEMBB100106	0 F-HEMBB1001060	1190	R-HEMBB1001060	6699
	HEMBB100106	3 F-HEMBB1001063	1191	R-HEMBB1001063	6700
	HEMBB100106	8 F-HEMBB1001068	1192	R-HEMBB1001068	6701
	HEMBB100109	6 F-HEMBB1001096	1193	R-HEMBB1001096	6702
35	HEMBB100110		1194	R-HEMBB1001102	6703
	HEMBB100110		1195	R-HEMBB1001105	6704
	HEMBB100111		1196		
	HEMBB100111		1197	R-HEMBB1001114	6705
40	HEMBB100111		1198	R-HEMBB1001117	6706
	HEMBB100111		1199	R-HEMBB1001119	6707
	HEMBB100112		1200	R-HEMBB1001126	6708
	HEMBB100113		1201	R-HEMBB1001133	6709
45	HEMBB100113		1202	R-HEMBB1001137	6710
	HEMBB100114		1203	R-HEMBB1001142	6711
	HEMBB100115		1204	R-HEMBB1001151	6712
	HEMBB100115		1205	R-HEMBB1001153	6713
50	HEMBB100116		1206	R-HEMBB1001169	6714
	HEMBB100117		1207	R-nnnnnnnnnnn	6715
	HEMBB100117		1208	R-HEMBB1001177	6716
	HEMBB100118		1209	R-HEMBB1001182	6717
55	HEMBB100119		1210	R-HEMBB1001199	6718
55	HEMBB100120	8 F-HEMBB1001208	1211	R-HEMBB1001208	6719

	HEMBB1001209	F-HEMBB1001209	1212	R-HEMBB1001209	6720
	HEMBB1001210	F-HEMBB1001210	1213	R-HEMBB1001210	6721
	HEMBB1001218	F-HEMBB1001218	1214	R-HEMBB1001218	6722
5	HEMBB1001221	F-HEMBB1001221	1215	R-HEMBB1001221	6723
	HEMBB1001234	F-HEMBB1001234	1216	R-HEMBB1001234	6724
	HEMBB1001242	F-HEMBB1001242	1217	R-HEMBB1001242	6725
	HEMBB1001249	F-HEMBB1001249	1218	R-HEMBB1001249	6726
10	HEMBB1001253	F-HEMBB1001253	1219	R-HEMBB1001253	6727
	HEMBB1001254	F-HEMBB1001254	1220	R-HEMBB1001254	6728
	HEMBB1001267	F-HEMBB1001267	1221	R-HEMBB1001267	6729
	HEMBB1001271	F-HEMBB1001271	1222	R-HEMBB1001271	6730
15	HEMBB1001282	F-HEMBB1001282	1223	R-HEMBB1001282	6731
	HEMBB1001288	F-HEMBB1001288	1224	R-HEMBB1001288	6732
	HEMBB1001289	F-HEMBB1001289,	1225	R-HEMBB1001289	6733
	HEMBB1001294	F-HEMBB1001294	1226	R-HEMBB1001294	6734
20	HEMBB1001302	F-HEMBB1001302	1227	R-HEMBB1001302	6735
	HEMBB1001304	F-HEMBB1001304	1228	R-HEMBB1001304	6736
	HEMBB1001314	F-HEMBB1001314	1229	R-HEMBB1001314	6737
	HEMBB1001315	F-HEMBB1001315	1230	R-HEMBB1001315	6738
25	HEMBB1001317	F-HEMBB1001317	1231	R-HEMBB1001317	6739
25	HEMBB1001326	F-HEMBB1001326	1232	R-HEMBB1001326	6740
	HEMBB1001331	F-HEMBB1001331	1233	R-HEMBB1001331	6741
	HEMBB1001335	F-HEMBB1001335	1234	R-HEMBB1001335	6742
•	HEMBB1001337	F-HEMBB1001337	1235	R-HEMBB1001337	6743
30	HEMBB1001339	F-HEMBB1001339	1236	R-HEMBB1001339	6744
	HEMBB1001346	F-HEMBB1001346	1237	R-HEMBB1001346	6745
	HEMBB1001348	F-HEMBB1001348	1238	R-HEMBB1001348	6746
	HEMBB1001356	F-HEMBB1001356	1239	R-HEMBB1001356	6747
35	HEMBB1001364	F-HEMBB1001364	1240	R-HEMBB1001364	6748
	HEMBB1001366	F-HEMBB1001366	1241	R-HEMBB1001366	6749
	HEMBB1001367	F-HEMBB1001367	1242	R-HEMBB1001367	6750
	HEMBB1001369	F-HEMBB1001369	1243	R-HEMBB1001369	6751
40	HEMBB1001380	F-HEMBB1001380	1244	R-HEMBB1001380	6752
	HEMBB1001384	F-HEMBB1001384	1245	R-HEMBB1001384	6753
	HEMBB1001387	F-HEMBB1001387	1246	R-HEMBB1001387	6754
	HEMBB1001394	F-HEMBB1001394	1247	R-HEMBB1001394	6755
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	HEMBB1001424	F-HEMBB1001424	1249	R-HEMBB1001424	6757
	HEMBB1001426	F-HEMBB1001426	1250	R-HEMBB1001426	6758
	HEMBB1001429	F-HEMBB1001429	1251	R-HEMBB1001429	6759
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	HEMBB1001443	F-HEMBB1001443	1253	R-HEMBB1001443	6761
	HEMBB1001449	F-HEMBB1001449	1254	R-HEMBB1001449	6762
	HEMBB1001454	F-HEMBB1001454	1255	R-HEMBB1001454	6763
55	HEMBB1001458	F-HEMBB1001458	1256	R-HEMBB1001458	6764
	HEMBB1001463	F-HEMBB1001463	1257	R-HEMBB1001463	6765

	HEMBB1001464	F-HEMBB1001464	1258	R-HEMBB1001464	6766
	HEMBB1001482	F-HEMBB1001482	1259	R-HEMBB1001482	6767
	HEMBB1001500	F-HEMBB1001500	1260	R-HEMBB1001500	6768
5	HEMBB1001521	F-HEMBB1001521	1261	R-HEMBB1001521	6769
	HEMBB1001527	F-HEMBB1001527	1262	R-HEMBB1001527	6770
	HEMBB1001531	F-HEMBB1001531	1263	R-HEMBB1001531	6771
	HEMBB1001535	F-HEMBB1001535	1264	R-HEMBB1001535	6772
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	HEMBB1001537	F-HEMBB1001537	1266	R-HEMBB1001537	6774
	HEMBB1001555	F-HEMBB1001555	1267	R-HEMBB1001555	6775
	HEMBB1001562	F-HEMBB1001562	1268	R-HEMBB1001562	6776
15	HEMBB1001564	F-HEMBB1001564	1269	R-HEMBB1001564	6777
	HEMBB1001565	F-HEMBB1001565	1270	R-HEMBB1001565	6778
	HEMBB1001585	F-HEMBB1001585,	1271	R-HEMBB1001585	6779
	HEMBB1001586	F-HEMBB1001586	1272	R-HEMBB1001586	6780
20	HEMBB1001588	F-HEMBB1001588	1273	R-HEMBB1001588	6781
20	HEMBB1001603	F-HEMBB1001603	1274	R-HEMBB1001603	6782
	HEMBB1001618	F-HEMBB1001618	1275	R-HEMBB1001618	6783
	HEMBB1001619	F-HEMBB1001619	1276	R-HEMBB1001619	6784
25	HEMBB1001630	F-HEMBB1001630	1277	R-HEMBB1001630	6785
25	HEMBB1001635	F-HEMBB1001635	1278	R-HEMBB1001635	6786
	HEMBB1001637	F-HEMBB1001637	1279	R-HEMBB1001637	6787
	HEMBB1001641	F-HEMBB1001641	1280	R-HEMBB1001641	6788
	HEMBB1001653	F-HEMBB1001653	1281	R-HEMBB1001653	6789
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	HEMBB1001668	F-HEMBB1001668	1283	R-HEMBB1001668	6791
	HEMBB1001673	F-HEMBB1001673	1284	R-HEMBB1001673	6792
	HEMBB1001684	F-HEMBB1001684	1285	R-HEMBB1001684	6793
35	HEMBB1001685	F-HEMBB1001685	1286	R-HEMBB1001685	6794
	HEMBB1001695	F-HEMBB1001695	1287	R-HEMBB1001695	6795
	HEMBB1001704	F-HEMBB1001704	1288	R-HEMBB1001704	6796
	HEMBB1001706	F-HEMBB1001706	1289	R-HEMBB1001706	6797
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	HEMBB1001717	F-HEMBB1001717	1291	R-HEMBB1001717	6799
	HEMBB1001735	F-HEMBB1001735	1292	R-HEMBB1001735	6800
	HEMBB1001736	F-HEMBB1001736	1293	R-HEMBB1001736	6801
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	HEMBB1001749	F-HEMBB1001749	1295	R-HEMBB1001749	6803
	HEMBB1001753	F-HEMBB1001753	1296	R-HEMBB1001753	6804
	HEMBB1001756	F-HEMBB1001756	1297	R-HEMBB1001756	6805
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	HEMBB1001785	F-HEMBB1001785	1300	R-HEMBB1001785	6808
	HEMBB1001797	F-HEMBB1001797	1301	R-HEMBB1001797	6809
66	HEMBB1001802	F-HEMBB1001802	1302	R-HEMBB1001802	6810
55	HEMBB1001812	F-HEMBB1001812	1303	R-HEMBB1001812	6811

	HEMBB1001816	F-HEMBB1001816	1304	R-HEMBB1001816	6812
	HEMBB1001831	F-HEMBB1001831	1305	R-HEMBB1001831	6813
	HEMBB1001834	F-HEMBB1001834	1306		
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	HEMBB1001839	F-HEMBB1001839	1308	R-HEMBB1001839	6815
	HEMBB1001850	F-HEMBB1001850	1309	R-HEMBB1001850	6816
	HEMBB1001863	F-HEMBB1001863	1310	R-HEMBB1001863	6817
10	HEMBB1001867	F-HEMBB1001867	1311	R-HEMBB1001867	6818
	HEMBB1001868	F-HEMBB1001868	1312	R-HEMBB1001868	6819
	HEMBB1001869	F-HEMBB1001869	1313	R-HEMBB1001869	6820
	HEMBB1001872	F-HEMBB1001872	1314	R-HEMBB1001872	6821
15	HEMBB1001874	F-HEMBB1001874	1315	R-HEMBB1001874	6822
	HEMBB1001875	F-HEMBB1001875	1316	R-HEMBB1001875	6823
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20	HEMBB1001905	F-HEMBB1001905	1319	R-HEMBB1001905	6826
20	HEMBB1001906	F-HEMBB1001906	1320	R-HEMBB1001906	6827
	HEMBB1001908	F-HEMBB1001908	1321	R-HEMBB1001908	6828
	HEMBB1001910	F-HEMBB1001910	1322	R-HEMBB1001910	6829
	HEMBB1001911	F-HEMBB1001911	1323	R-HEMBB1001911	6830
25	HEMBB1001915	F-HEMBB1001915	1324	R-HEMBB1001915	6831
	HEMBB1001921	F-HEMBB1001921	1325	R-HEMBB1001921	6832
	HEMBB1001922	F-HEMBB1001922	1326	R-HEMBB1001922	6833
	HEMBB1001925	F-HEMBB1001925	1327	R-HEMBB1001925	6834
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	HEMBB1001944	F-HEMBB1001944	1329	R-HEMBB1001944	6836
	HEMBB1001945	F-HEMBB1001945	1330	R-HEMBB1001945	6837
	HEMBB1001947	F-HEMBB1001947	1331	R-HEMBB1001947	6838
35	HEMBB1001950	F-HEMBB1001950	1332	R-HEMBB1001950	6839
	HEMBB1001952	F-HEMBB1001952	1333	R-HEMBB1001952	6840
	HEMBB1001953	F-HEMBB1001953	1334	R-HEMBB1001953	6841
	HEMBB1001957	F-HEMBB1001957	1335	R-HEMBB1001957	6842
40	HEMBB1001962	F-HEMBB1001962	1336	R-HEMBB1001962	6843
	HEMBB1001967	F-HEMBB1001967	1337	R-HEMBB1001967	6844
	HEMBB1001973	F-HEMBB1001973	1338	R-HEMBB1001973	6845
	HEMBB1001983	F-HEMBB1001983	1339	R-HEMBB1001983	6846
45	HEMBB1001988	F-HEMBB1001988	1340	R-HEMBB1001988	6847
	HEMBB1001990	F-HEMBB1001990	1341	R-HEMBB1001990	6848
	HEMBB1001996	F-HEMBB1001996	1342	R-HEMBB1001996	6849
	HEMBB1001997	F-HEMBB1001997	1343	R-HEMBB1001997	6850
50	HEMBB1002002	F-HEMBB1002002	1344	R-HEMBB1002002	6851
	HEMBB1002005	F-HEMBB1002005	1345	R-HEMBB1002005	6852
	HEMBB1002009	F-HEMBB1002009	1346	R-HEMBB1002009	6853
	HEMBB1002015	F-HEMBB1002015	1347	R-HEMBB1002015	6854
£ £	HEMBB1002042	F-HEMBB1002042	1348	R-HEMBB1002042	6855
55	HEMBB1002043	F-HEMBB1002043	1349	R-HEMBB1002043	6856

	HEMBB1002044	F-HEMBB1002044	1350	R-HEMBB1002044	6857
	HEMBB1002045	F-HEMBB1002045	1351	R-HEMBB1002045	6858
	HEMBB1002049	F-HEMBB1002049	1352	R-HEMBB1002049	6859
5	HEMBB1002050	F-HEMBB1002050	1353	R-HEMBB1002050	6860
	HEMBB1002068	F-HEMBB1002068	1354	R-HEMBB1002068	6861
	HEMBB1002069	F-HEMBB1002069	1355	R-HEMBB1002069	6862
	HEMBB1002092	F-HEMBB1002092	1356	R-HEMBB1002092	6863
10	HEMBB1002094	F-HEMBB1002094	1357	R-HEMBB1002094	6864
	HEMBB1002115	F-HEMBB1002115	1358	R-HEMBB1002115	6865
	HEMBB1002134	F-HEMBB1002134	1359		
	HEMBB1002139	F-HEMBB1002139	1360	R-HEMBB1002139	6866
15	HEMBB1002142	F-HEMBB1002142	1361	R-HEMBB1002142	6867
, ,	HEMBB1002152	F-HEMBB1002152	1362	R-HEMBB1002152	6868
	HEMBB1002189	F-HEMBB1002189,	1363	R-HEMBB1002189	6869
	HEMBB1002190	F-HEMBB1002190	1364	R-HEMBB1002190	6870
20	HEMBB1002193	F-HEMBB1002193	1365	R-HEMBB1002193	6871
20	HEMBB1002217	F-HEMBB1002217	1366	R-HEMBB1002217	6872
	HEMBB1002218	F-HEMBB1002218	1367	R-HEMBB1002218	6873
	HEMBB1002232	F-HEMBB1002232	1368	R-HEMBB1002232	6874
9.5	HEMBB1002247	F-HEMBB1002247	1369	R-HEMBB1002247	6875
25	HEMBB1002249	F-HEMBB1002249	1370	R-HEMBB1002249	6876
	HEMBB1002254	F-HEMBB1002254	1371	R-HEMBB1002254	6877
	HEMBB1002255	F-HEMBB1002255	1372	R-HEMBB1002255	6878
	HEMBB1002266	F-HEMBB1002266	1373	R-HEMBB1002266	6879
30	HEMBB1002280	F-HEMBB1002280	1374	R-HEMBB1002280	6880
	HEMBB1002300	F-HEMBB1002300	1375	R-HEMBB1002300	6881
	HEMBB1002306	F-HEMBB1002306	1376	R-HEMBB1002306	6882
	HEMBB1002327	F-HEMBB1002327	1377	R-HEMBB1002327	6883
35	HEMBB1002329	F-HEMBB1002329	1378	R-HEMBB1002329	6884
	HEMBB1002340	F-HEMBB1002340	1379	R-HEMBB1002340	6885
	HEMBB1002342	F-HEMBB1002342	1380	R-HEMBB1002342	6886 6887
	HEMBB1002358	F-HEMBB1002358	1381	R-HEMBB1002358	6888
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	HEMBB1002364	F-HEMBB1002364	1383		6890
	HEMBB1002371	F-HEMBB1002371 F-HEMBB1002381	1384 1385	R-HEMBB1002371 R-HEMBB1002381	6891
	HEMBB1002381 HEMBB1002383	F-HEMBB1002383	1386	R-HEMBB1002383	6892
45	HEMBB1002387	F-HEMBB1002387	1387	R-HEMBB1002387	6893
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	HEMBB1002415	F-HEMBB1002425	1390	R-HEMBB1002425	6895
50	HEMBB1002442	F-HEMBB1002442	1391	R-HEMBB1002442	6896
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	HEMBB1002457	F-HEMBB1002457	1393	R-HEMBB1002457	6898
	HEMBB1002458		1394	R-HEMBB1002458	6899
55	HEMBB1002477		1395	R-HEMBB1002477	6900

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	HEMBB1002492	F-HEMBB1002492	1397	R-HEMBB1002492	6902
	HEMBB1002495	F-HEMBB1002495	1398	R-HEMBB1002495	6903
5	HEMBB1002502	F-HEMBB1002502	1399	R-HEMBB1002502	6904
	HEMBB1002509	F-HEMBB1002509	1400	R-HEMBB1002509	6905
	HEMBB1002510	F-HEMBB1002510	1401	R-HEMBB1002510	6906
	HEMBB1002520	F-HEMBB1002520	1402	R-HEMBB1002520	6907
10	HEMBB1002522	F-HEMBB1002522	1403	R-HEMBB1002522	6908
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	HEMBB1002534	F-HEMBB1002534	1405	R-HEMBB1002534	6910
	HEMBB1002545	F-HEMBB1002545	1406	R-HEMBB1002545	6911
15	HEMBB1002550	F-HEMBB1002550	1407	R-HEMBB1002550	6912
	HEMBB1002556	F-HEMBB1002556	1408	R-HEMBB1002556	6913
	HEMBB1002579	F-HEMBB1002579,		R-HEMBB1002579	6914
	HEMBB1002582	F-HEMBB1002582	1410	R-HEMBB1002582	6915
20	HEMBB1002590	F-HEMBB1002590	1411	R-HEMBB1002590	6916
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	HEMBB1002600	F-HEMBB1002600	1413	R-HEMBB1002600	6918
	HEMBB1002601	F-HEMBB1002601	1414	R-HEMBB1002601	6919
	HEMBB1002603	F-HEMBB1002603	1415	R-HEMBB1002603	6920
25	HEMBB1002607	F-HEMBB1002607	1416	R-HEMBB1002607	6921
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	HEMBB1002613	F-HEMBB1002613	1418	R-HEMBB1002613	6923
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	HEMBB1002623	F-HEMBB1002623	1421	R-HEMBB1002623	6926
	HEMBB1002635	F-HEMBB1002635	1422	R-HEMBB1002635	6927
	HEMBB1002664	F-HEMBB1002664	1423	R-HEMBB1002664	6928
35	HEMBB1002677	F~HEMBB1002677	1424	R-HEMBB1002677	6929
	HEMBB1002683	F-HEMBB1002683	1425	R-HEMBB1002683	6930
	HEMBB1002684	F-HEMBB1002684	1426	R-HEMBB1002684	6931
	HEMBB1002686	F-HEMBB1002686	1427	R-HEMBB1002686	6932
40	HEMBB1002692	F-HEMBB1002692	1428	R-HEMBB1002692	6933
	HEMBB1002697	F-HEMBB1002697	1429	R-HEMBB1002697	6934
	HEMBB1002699	F-HEMBB1002699	1430	R-HEMBB1002699	6935
	HEMBB1002702	F-HEMBB1002702	1431	R-HEMBB1002702	6936
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45	HEMBB1002712	F-HEMBB1002712	1433	R-HEMBB1002712	6938
	MAMMA1000009	F-MAMMA1000009	1434	R-MAMMA1000009	6939
	MAMMA1000019	F-MAMMA1000019	1435	R-MAMMA1000019	6940
	MAMMA1000020	F-MAMMA1000020	1436	R-MAMMA1000020	6941
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	MAMMA1000043	F-MAMMA1000043	1438	R-MAMMA1000043	6943
	MAMMA1000045	F-MAMMA1000045	1439	R-MAMMA1000045	6944
	MAMMA1000055	F-MAMMA1000055	1440	R-MAMMA1000055	6945
55	MAMMA1000057	F-MAMMA1000057	1441	R-MAMMA1000057	6946

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	MAMMA10	00085 F-MAMMA	1000085 1444	R-MAMMA1000085	6949
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	MAMMA10	00103 F-MAMMA	1000103 1446	R-MAMMA1000103	6951
	MAMMA10		1000117 1447		6952
	MAMMA10		1000129 1448	R-MAMMA1000129	6953
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	MAMMA10		1000134 1450		6955
	MAMMA10	00139 F-MAMM	1000139 1451	R-MAMMA1000139	6956
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15	MAMMA10		1000155 1453		6958
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	MAMMA10		1000173 1456	6 R-MAMMA1000173	6961
20	MAMMA10	00175 F-MAMM	1000175 1457	7 R-MAMMA1000175	6962
20	MAMMA10	00183 F-MAMM/	1000183 1458	R-MAMMA1000183	6963
	MAMMA10	00198 F-MAMM	11000198 1459	R-MAMMA1000198	6964
	MAMMA10	00221 F-MAMM	1000221 1460	R-MAMMA1000221	6965
	MAMMA10	00227 F-MAMM	1000227 146	R-MAMMA1000227	6966
25	MAMMA10	00241 F-MAMM	1000241 1462	R-MAMMA1000241	6967
	MAMMA10	00251 F-MAMM	11000251 1463	R-MAMMA1000251	
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40	MAMMA10		11000302 1474		
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	MAMMA10		11000309 1470		
	MAMMA10		11000312 147		
45	MAMMA10		11000313 1478		
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	MAMMA10		1000339 1480		
	MAMMA10		A1000340 1483 A1000348 1483		
50	MAMMA10 MAMMA10		\1000348		
	MAMMA10 MAMMA10		A1000360 1484 A1000361 148		
			A1000361 1489 A1000372 1489	· · · · · · · · · · · · · · · · · · ·	
55	MAMMA10				
55	MAMMA10	UU385 F-MAMM	A1000385 148'	1 V_WVWWYTOOOS	0334

	- MAMMA1000388	F-MAMMA1000388	1488	R-MAMMA1000388	6993
	MAMMA1000395	F-MAMMA1000395	1489	R-MAMMA1000395	6994
	MAMMA1000402	F-MAMMA1000402	1490	R-MAMMA1000402	6995
5	MAMMA1000410	F-MAMMA1000410	1491	R-MAMMA1000410	6996
	MAMMA1000413	F-MAMMA1000413	1492	R-MAMMA1000413	6997
	MAMMA1000414	F-MAMMA1000414	1493	R-MAMMA1000414	6998
	MAMMA1000416	F-MAMMA1000416	1494	R-MAMMA1000416	6999
10	MAMMA1000421	F-MAMMA1000421	1495	R-MAMMA1000421	7000
	MAMMA1000422	F-MAMMA1000422	1496	R-MAMMA1000422	7001
	MAMMA1000423	F-MAMMA1000423	1497	R-MAMMA1000423	7002
	MAMMA1000424	F-MAMMA1000424	1498	R-MAMMA1000424	7003
15	MAMMA1000429	F-MAMMA1000429	1499	R-MAMMA1000429	7004
	MAMMA1000431	F-MAMMA1000431	1500	R-MAMMA1000431	7005
	MAMMA1000444	F-MAMMA1000444,	1501	R-MAMMA1000444	7006
	MAMMA1000446	F-MAMMA1000446	1502	R-MAMMA1000446	7007
20	MAMMA1000458	F-MAMMA1000458	1503	R-MAMMA1000458	7008
20	MAMMA1000468	F-MAMMA1000468	1504	R-MAMMA1000468	7009
	MAMMA1000472	F-MAMMA1000472	1505	R-MAMMA1000472	7010
	MAMMA1000478	F-MAMMA1000478	1506	R-MAMMA1000478	7011
	MAMMA1000483	F-MAMMA1000483	1507	R-MAMMA1000483	7012
25	MAMMA1000490	F-MAMMA1000490	1508	R-MAMMA1000490	7013
	MAMMA1000500	F-MAMMA1000500	1509	R-MAMMA1000500	7014
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	MAMMA1000516	F-MAMMA1000516	1511	R-MAMMA1000516	7016
30	MAMMA1000522	F-MAMMA1000522	1512	R-MAMMA1000522	7017
	MAMMA1000524	F-MAMMA1000524	1513		
	MAMMA1000559	F-MAMMA1000559	1514	R-MAMMA1000559	7018
	MAMMA1000565	F-MAMMA1000565	1515	R-MAMMA1000565	7019
35	MAMMA1000567	F-MAMMA1000567	1516	R-MAMMA1000567	7020
	MAMMA1000576	F-MAMMA1000576	1517	R-MAMMA1000576	7021
	MAMMA1000583	F-MAMMA1000583	1518	R-MAMMA1000583	7022
	MAMMA1000585	F-MAMMA1000585	1519	R-MAMMA1000585	7023
40	MAMMA1000594	F-MAMMA1000594	1520	R-MAMMA1000594	7024
	MAMMA1000597	F-MAMMA1000597	1521	R-MAMMA1000597	7025
	MAMMA1000605	F-MAMMA1000605	1522	R-MAMMA1000605	7026
	MAMMA1000612	F-MAMMA1000612	1523	R-MAMMA1000612	7027
45	MAMMA1000616	F-MAMMA1000616	1524	R-MAMMA1000616	7028
	MAMMA1000621	F-MAMMA1000621	1525	R-MAMMA1000621	7029
	MAMMA1000623	F-MAMMA1000623	1526	R-MAMMA1000623	7030
	MAMMA1000625	F-MAMMA1000625	1527	R-MAMMA1000625	7031
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	MAMMA1000669	F-MAMMA1000669	1530	R-MAMMA1000669	7034
	MAMMA1000670	F-MAMMA1000670	1531	R-MAMMA1000670	7035
	MAMMA1000672	F-MAMMA1000672	1532	R-MAMMA1000672	7036
55	MAMMA1000684	F-MAMMA1000684	1533	R-MAMMA1000684	7037

	MAMMA1000696	F-MAMMA1000696	1534	R-MAMMA1000696	7038
	MAMMA1000707	F-MAMMA1000707	1535	R-MAMMA1000707	7039
	MAMMA1000713	F-MAMMA1000713	1536	R-MAMMA1000713	7040
5	MAMMA1000714	F-MAMMA1000714	1537	R-MAMMA1000714	7041
	MAMMA1000718	F-MAMMA1000718	1538	R-MAMMA1000718	7042
	MAMMA1000720	F-MAMMA1000720	1539	R-MAMMA1000720	7043
	MAMMA1000723	F-MAMMA1000723	1540	R-MAMMA1000723	7044
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	MAMMA1000732	F-MAMMA1000732	1542	R-MAMMA1000732	7046
	MAMMA1000733	F-MAMMA1000733	1543	R-MAMMA1000733	7047
	MAMMA1000734	F-MAMMA1000734	1544	R-MAMMA1000734	7048
15	MAMMA1000738	F-MAMMA1000738	1545	R-MAMMA1000738	7049
	MAMMA1000744	F-MAMMA1000744	1546	R-MAMMA1000744	7050
	MAMMA1000746	F-MAMMA1000746,	1547	R-MAMMA1000746	7051
	MAMMA1000752	F-MAMMA1000752	1548	R-MAMMA1000752	7052
20	MAMMA1000760	F-MAMMA1000760	1549	R-MAMMA1000760	7053
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	MAMMA1000775	F-MAMMA1000775	1551	R-MAMMA1000775	7055
	MAMMA1000776	F-MAMMA1000776	1552	R-MAMMA1000776	7056
	MAMMA1000778	F-MAMMA1000778	1553	R-MAMMA1000778	7057
25	MAMMA1000782	F-MAMMA1000782	1554	R-MAMMA1000782	7058
	MAMMA1000798	F-MAMMA1000798	1555	R-MAMMA1000798	7059
	MAMMA1000802	F-MAMMA1000802	1556	R-MAMMA1000802	7060
	MAMMA1000824	F-MAMMA1000824	1557		
30	MAMMA1000831	F-MAMMA1000831	1558	R-MAMMA1000831	7061
	MAMMA1000839	F-MAMMA1000839	1559	R-MAMMA1000839	7062
	MAMMA1000841	F-MAMMA1000841	1560	R-MAMMA1000841	7063
	MAMMA1000842	F-MAMMA1000842	1561	R-MAMMA1000842	7064
35	MAMMA1000843	F-MAMMA1000843	1562	R-MAMMA1000843	7065
	MAMMA1000845	F-MAMMA1000845	1563	R-MAMMA1000845	7066
	MAMMA1000851	F-MAMMA1000851	1564	R-MAMMA1000851	7067
	MAMMA1000855	F-MAMMA1000855	1565	R-MAMMA1000855	7068
40	MAMMA1000856	F-MAMMA1000856	1566	R-MAMMA1000856	7069
	MAMMA1000859	F-MAMMA1000859	1567		
	MAMMA1000862	F-MAMMA1000862	1568	R-MAMMA1000862	7070
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40	MAMMA1000867	F-MAMMA1000867	1571	R-MAMMA1000867	7073
	MAMMA1000875	F-MAMMA1000875	1572	R-MAMMA1000875	7074
	MAMMA1000876	F-MAMMA1000876	1573	R-MAMMA1000876	7075
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50	MAMMA1000880	F-MAMMA1000880	1575	R-MAMMA1000880	7077
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	MAMMA1000897	F-MAMMA1000897	1577	R-MAMMA1000897	7079
	MAMMA1000905	F-MAMMA1000905	1578	R-MAMMA1000905	7080
55	MAMMA1000906	F-MAMMA1000906	1579	R-MAMMA1000906	7081

	MAMMA1000908	F-MAMMA1000908	1580	R-MAMMA1000908	7082
	MAMMA1000914	F-MAMMA1000914	1581	R-MAMMA1000914	7083
	MAMMA1000921	F-MAMMA1000921	1582	R-MAMMA1000921	7084
5	MAMMA1000931	F-MAMMA1000931	1583	R-MAMMA1000931	7085
	MAMMA1000940	. F-MAMMA1000940	1584	R-MAMMA1000940	7086
	MAMMA1000941	F-MAMMA1000941	1585	R-MAMMA1000941	7087
	MAMMA1000942	F-MAMMA1000942	1586	R-MAMMA1000942	7088
10	MAMMA1000943	F-MAMMA1000943	1587	R-MAMMA1000943	7089
	MAMMA1000956	F-MAMMA1000956	1588	R-MAMMA1000956	7090
	MAMMA1000957	F-MAMMA1000957	1589	R-MAMMA1000957	7091
	MAMMA1000962	F-MAMMA1000962	1590	R-MAMMA1000962	7092
15	MAMMA1000968	F-MAMMA1000968	1591	R-MAMMA1000968	7093
	MAMMA1000975	F-MAMMA1000975	1592	R-MAMMA1000975	7094
	MAMMA1000979	F-MAMMA1000979,	1593	R-MAMMA1000979	7095
	MAMMA1000987	F-MAMMA1000987	1594	R-MAMMA1000987	7096
20	MAMMA1000998	F-MAMMA1000998	1595	R-MAMMA1000998	7097
20	MAMMA1001003	F-MAMMA1001003	1596	R-MAMMA1001003	7098
	MAMMA1001008	F-MAMMA1001008	1597	R-MAMMA1001008	7099
	MAMMA1001021	F-MAMMA1001021	1598	R-MAMMA1001021	7100
	MAMMA1001024	F-MAMMA1001024	1599	R-MAMMA1001024	7101
25	MAMMA1001030	F-MAMMA1001030	1600	R-MAMMA1001030	7102
	MAMMA1001035	F-MAMMA1001035	1601	R-MAMMA1001035	7103
	MAMMA1001038	F-MAMMA1001038	1602	R-MAMMA1001038	7104
	MAMMA1001041	F-MAMMA1001041	1603	R-nnnnnnnnnn	7105
30	MAMMA1001050	F-MAMMA1001050	1604	R-MAMMA1001050	7106
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	MAMMA1001067	F-MAMMA1001067	1606	R-MAMMA1001067	7108
	MAMMA1001073	F-MAMMA1001073	1607	R-MAMMA1001073	7109
35	MAMMA1001074	F-MAMMA1001074	1608	R-MAMMA1001074	7110
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	MAMMA1001078	F-MAMMA1001078	1610	R-MAMMA1001078	7112
	MAMMA1001080	F-MAMMA1001080	1611		
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	MAMMA1001091	F-MAMMA1001091	1613	R-MAMMA1001091	7114
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45	MAMMA1001110	F-MAMMA1001110	1616	R-MAMMA1001110	7117
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	MAMMA1001133	F-MAMMA1001133	1618	R-MAMMA1001133	7119
	MAMMA1001139	F-MAMMA1001139	1619	R-MAMMA1001139	7120
50	MAMMA1001143	F-MAMMA1001143	1620	R-MAMMA1001143	7121
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	MAMMA1001154	F-MAMMA1001154	1622	R-MAMMA1001154	7123
	MAMMA1001161	F-MAMMA1001161	1623	R-MAMMA1001161	7124
	MAMMA1001162	F-MAMMA1001162	1624	R-MAMMA1001162	7125
55	MAMMA1001181	F-MAMMA1001181	1625	R-MAMMA1001181	7126

	MAMMA1001186	F-MAMMA1001186	1626	R-MAMMA1001186	7127
	MAMMA1001191		1627	R-MAMMA1001191	7128
		F-MAMMA1001191			
5	MAMMA1001198	F-MAMMA1001198	1628	R-MAMMA1001198	7129
	MAMMA1001202	F-MAMMA1001202	1629	R-MAMMA1001202	7130
	MAMMA1001203	F-MAMMA1001203	1630	R-MAMMA1001203	7131
	MAMMA1001206	F-MAMMA1001206	1631	R-MAMMA1001206	7132
	MAMMA1001215	F-MAMMA1001215	1632	R-MAMMA1001215	7133
10	MAMMA1001220	F-MAMMA1001220	1633	R-MAMMA1001220	7134
	MAMMA1001222	F-MAMMA1001222	1634	R-MAMMA1001222	7135
	MAMMA1001243	F-MAMMA1001243	1635	R-MAMMA1001243	7136
	MAMMA1001244	F-MAMMA1001244	1636	R-MAMMA1001244	7137
15	MAMMA1001249	F-MAMMA1001249	1637	R-MAMMA1001249	7138
	MAMMA1001256	F-MAMMA1001256	1638	R-MAMMA1001256	7139
	MAMMA1001259	F-MAMMA1001259	1639	R-MAMMA1001259	7140
	MAMMA1001260	F-MAMMA1001260	1640	R-MAMMA1001260	7141
20	MAMMA1001268	F-MAMMA1001268	1641	R-MAMMA1001268	7142
	MAMMA1001271	F-MAMMA1001271	1642	R-MAMMA1001271	7143
	MAMMA1001274	F-MAMMA1001274	1643	R-MAMMA1001274	7144
	MAMMA1001280	F-MAMMA1001280	1644	R-MAMMA1001280	7145
0.5	MAMMA1001292	F-MAMMA1001292	1645	R-MAMMA1001292	7146
25	MAMMA1001296	F-MAMMA1001296	1646	R-MAMMA1001296	7147
	MAMMA1001298	F-MAMMA1001298	1647	R-MAMMA1001298	7148
	MAMMA1001305	F-MAMMA1001305	1648	R-MAMMA1001305	7149
	MAMMA1001322	F-MAMMA1001322	1649	R-MAMMA1001322	7150
30	MAMMA1001324	F-MAMMA1001324	1650	R-MAMMA1001324	7151
	MAMMA1001330	F-MAMMA1001330	1651	R-MAMMA1001330	7152
	MAMMA1001341	F-MAMMA1001341	1652	R-MAMMA1001341	7153
	MAMMA1001343	F-MAMMA1001343	1653	R-MAMMA1001343	7154
35	MAMMA1001346	F-MAMMA1001346	1654	R-MAMMA1001346	7155
	MAMMA1001383	F-MAMMA1001383	1655	R-MAMMA1001383	7156
	MAMMA1001388	F-MAMMA1001388	1656	R-MAMMA1001388	7157
	MAMMA1001397	F-MAMMA1001397	1657	R-MAMMA1001397	7158
40	MAMMA1001408	F-MAMMA1001408	1658	R-MAMMA1001408	7159
40	MAMMA1001411	F-MAMMA1001411	1659	R-MAMMA1001411	7160
	MAMMA1001419	F-MAMMA1001419	1660	R-MAMMA1001419	7161
	MAMMA1001420	F-MAMMA1001420	1661	R-MAMMA1001420	7162
	MAMMA1001435	F-MAMMA1001435	1662	R-MAMMA1001435	7163
45	MAMMA1001442	F-MAMMA1001442	1663	R-MAMMA1001442	7164
	MAMMA1001446	F-MAMMA1001446	1664	R-MAMMA1001446	7165
	MAMMA1001452	F-MAMMA1001452	1665	R-MAMMA1001452	7166
	MAMMA1001465	F-MAMMA1001465	1666	R-MAMMA1001465	7167
50	MAMMA1001476	F-MAMMA1001476	1667	R-MAMMA1001476	7168
	MAMMA1001487	F-MAMMA1001487	1668	R-MAMMA1001487	7169
	MAMMA1001501	F-MAMMA1001501	1669	R-MAMMA1001501	7170
	MAMMA1001502	F-MAMMA1001502	1670	R-MAMMA1001502	7171
55	MAMMA1001510	F-MAMMA1001510	1671	R-MAMMA1001510	7172

	MAMMA1001522	F-MAMMA1001522	1672	R-MAMMA1001522	7173
	MAMMA1001547	F-MAMMA1001547	1673	R-MAMMA1001547	7174
	MAMMA1001551	F-MAMMA1001551	1674	R-MAMMA1001551	7175
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	MAMMA1001576	F-MAMMA1001576	1676	R-MAMMA1001576	7177
	MAMMA1001590	F-MAMMA1001590	1677	R-MAMMA1001590	7178
	MAMMA1001600	F-MAMMA1001600	1678	R-MAMMA1001600	7179
10	MAMMA1001604	F-MAMMA1001604	1679	R-MAMMA1001604	7180
	MAMMA1001606	F-MAMMA1001606	1680	R-MAMMA1001606	7181
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	MAMMA1001627	F-MAMMA1001627	1682	R-MAMMA1001627	7183
15	MAMMA1001630	F-MAMMA1001630	1683	R-MAMMA1001630	7184
,,	MAMMA1001633	F-MAMMA1001633	1684	R-MAMMA1001633	7185
	MAMMA1001635	F-MAMMA1001635,	1685	R-MAMMA1001635	7186
	MAMMA1001649	F-MAMMA1001649	1686	R-MAMMA1001649	7187
	MAMMA1001654	F-MAMMA1001654	1687		
20	MAMMA1001663	F-MAMMA1001663	1688	R-MAMMA1001663	7188
	MAMMA1001670	F-MAMMA1001670	1689	R-MAMMA1001670	7189
	MAMMA1001671	F-MAMMA1001671	1690	R-MAMMA1001671	7190
	MAMMA1001679	F-MAMMA1001679	1691	R-MAMMA1001679	7191
25	MAMMA1001683	F-MAMMA1001683	1692	R-MAMMA1001683	7192
	MAMMA1001686	F-MAMMA1001686	1693	R-MAMMA1001686	7193
	MAMMA1001692	F-MAMMA1001692	1694	R-MAMMA1001692	7194
	MAMMA1001711	F-MAMMA1001711	1695	R-MAMMA1001711	7195
30	MAMMA1001715	F-MAMMA1001715	1696	R-MAMMA1001715	7196
	MAMMA1001730	F-MAMMA1001730	1697	R-MAMMA1001730	7197
	MAMMA1001735	F-MAMMA1001735	1698	R-MAMMA1001735	7198
	MAMMA1001740	F-MAMMA1001740	1699	R-MAMMA1001740	7199
35	MAMMA1001743	F-MAMMA1001743	1700	R-MAMMA1001743	7200
	MAMMA1001744	F-MAMMA1001744	1701	R-MAMMA1001744	7201
	MAMMA1001745	F-MAMMA1001745	1702	R-MAMMA1001745	7202
	MAMMA1001751	F-MAMMA1001751	1703	R-MAMMA1001751	7203
40	MAMMA1001754	F-MAMMA1001754	1704	R-MAMMA1001754	7204
	MAMMA1001757	F-MAMMA1001757	1705	R-MAMMA1001757	7205
	MAMMA1001760	F-MAMMA1001760	1706	R-MAMMA1001760	7206
	MAMMA1001764	F-MAMMA1001764	1707	R-MAMMA1001764	7207
45	MAMMA1001768	F-MAMMA1001768	1708	R-MAMMA1001768	7208
45	MAMMA1001769	F-MAMMA1001769	1709	R-MAMMA1001769	7209
	MAMMA1001771	F-MAMMA1001771	1710	R-MAMMA1001771	7210
	MAMMA1001783	F-MAMMA1001783	1711	R-MAMMA1001783	7211
	MAMMA1001785	F-MAMMA1001785	1712	R-MAMMA1001785	7212
50	MAMMA1001788	F-MAMMA1001788	1713	R-MAMMA1001788	7213
	MAMMA1001790	F-MAMMA1001790	1714	R-MAMMA1001790	7214
	MAMMA1001806	F-MAMMA1001806	1715	R-MAMMA1001806	7215
	MAMMA1001812	F-MAMMA1001812	1716	R-MAMMA1001812	7216
55	MAMMA1001815	F-MAMMA1001815	1717	R-MAMMA1001815	7217

	MAMMA100181	7 F-MAMMA1001817	1718	R-MAMMA1001817	7218
	MAMMA100181	8 F-MAMMA1001818	1719	R-MAMMA1001818	7219
	MAMMA100182	O F-MAMMA1001820	1720	R-MAMMA1001820	7220
5	MAMMA100182			R-MAMMA1001824	7221
	MAMMA100183			R-MAMMA1001836	7222
	MAMMA100183		1723	R-MAMMA1001837	7223
	MAMMA100184		1724	R-MAMMA1001848	7224
10	MAMMA100185		1725	R-MAMMA1001851	7225
	MAMMA100185			R-MAMMA1001854	7226
	MAMMA100185		1727	R-MAMMA1001858	7227
	MAMMA100186		1728	R-MAMMA1001864	7228
	MAMMA100186				
15			1729	R-nnnnnnnnnnnn	7229
	MAMMA100187		1730	R-MAMMA1001874	7230
	MAMMA100187		•	R-MAMMA1001878	7231
	MAMMA100188	· ·	1732	R-MAMMA1001880	7232
20	MAMMA100189		1733	R-MAMMA1001890	7233
	MAMMA100190		1734	R-MAMMA1001907	7234
	MAMMA100190		1735	R-nnnnnnnnnnn	7235
	MAMMA100193		1736	R-MAMMA1001931	7236
25	MAMMA100195		1737	R-MAMMA1001956	7237
	MAMMA100196		1738	R-MAMMA1001963	7238
	MAMMA1001969		1739	R-MAMMA1001969	7239
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	MAMMA1002145		1762	R-MAMMA1002145	7262
55	MAMMA1002153		1763	R-MAMMA1002153	7263

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	MAMMA1002156	F-MAMMA1002156	1765	R-MAMMA1002156	7265
	MAMMA1002158	F-MAMMA1002158	1766	R-MAMMA1002158	7266
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	MAMMA1002230	F-MAMMA1002230	1773	R-MAMMA1002230	7273
	MAMMA1002236	F-MAMMA1002236	1774	R-MAMMA1002236	7274
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	MAMMA1002417	F-MAMMA1002427	1815	R-MAMMA1002427	7315
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	MAMMA1002461	F-MAMMA1002461	1820	R-MAMMA1002461	7320
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	MAMMA1002530	F-MAMMA1002530	1828	R-MAMMA1002530	7328
	MAMMA1002545	F-MAMMA1002545	1829	R-MAMMA1002545	7329
25	MAMMA1002554	F-MAMMA1002554	1830	R-MAMMA1002554	7330
	MAMMA1002556	F-MAMMA1002556	1831	R-MAMMA1002556	7331
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	MAMMA1002571	F-MAMMA1002571	1833	R-MAMMA1002571	7333
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	MAMMA1002590	F-MAMMA1002590	1836	R-MAMMA1002590	7336
	MAMMA1002597	F-MAMMA1002597	1837	R-MAMMA1002597	7337
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35	MAMMA1002603	F-MAMMA1002603	1839	R-MAMMA1002603	7339
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	MAMMA1002754	F-MAMMA1002754	1870	R-MAMMA1002754	7370
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55	1117/MIZOO1910	I INI ZIUUZOVIO+U	2101

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	NT2RM4000202	F-NT2RM4000202	2233	R-NT2RM4000202	7455
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30	NT2RM4000366	F-NT2RM4000366	2248	R-NT2RM4000366	7470
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	NT2RM4001256	F-NT2RM4001256	2321	R-NT2RM4001256	7543
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	NT2RM4001754	F-NT2RM4001754	2366	R-NT2RM4001754	7588
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	NT2RM4002073 NT2RM4002075	F-NT2RM4002073 F-NT2RM4002075	2404 2405	R-NT2RM4002075	7625
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	NT2RP2000503	F-NT2RP2000503	2635	R-NT2RP2000503	7729
55	NT2RP2000510	F-NT2RP2000510	2636	R-NT2RP2000510	7730
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	NT2RP2000603	F-NT2RP2000603	2639	R-NT2RP2000603	7732
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	NT2RP2000644	F-NT2RP2000644	2642	R-NT2RP2000644	7735
	NT2RP2000656	F-NT2RP2000656	2643	R-NT2RP2000656	7736
	NT2RP2000658	F-NT2RP2000658	2644	R-NT2RP2000658	7737
10	NT2RP2000668	F-NT2RP2000668	2645	R-NT2RP2000668	7738
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	NT2RP2000758				
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	NT2RP2000841	F-NT2RP2000841	2658	R-NT2RP2000841	7750
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	NT2RP2000892	F-NT2RP2000892	2663	R-NT2RP2000892	7755
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	NT2RP2000943	F-NT2RP2000943	2667	R-NT2RP2000943	7758
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	NT2RP2001036	F-NT2RP2001036	2672	R-NT2RP2001036	7763
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	NT2RP2001070	F-NT2RP2001070	2676	R-NT2RP2001070	7766
	NT2RP2001081	F-NT2RP2001081	2677		
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55	NT2RP2001168	F-NT2RP2001168	2683	R-NT2RP2001168	7772
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	NT2RP2001394	F-NT2RP2001394	2703	R-NT2RP2001394	7791
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	NT2RP2001721	F-NT2RP2001721	2735	R-NT2RP2001721	7820
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	NT2RP2001883	F-NT2RP2001883	2744	R-NT2RP2001883	7828
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	NT2RP2001969	F-NT2RP2001969	2753	R-NT2RP2001969	7836
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	NT2RP2001991	F-NT2RP2001991	2756		
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	NT2RP2002033	F-NT2RP2002033	2759	R-NT2RP2002033	7841
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55	NT2RP2002185	F-NT2RP2002185	2775	R-NT2RP2002185	7856

	NT2RP2002192	F-NT2RP2002192	2776	R-NT2RP2002192	7857
	NT2RP2002193	F-NT2RP2002193	2777	R-NT2RP2002193	7858
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	NT2RP2002235	F-NT2RP2002235	2781		
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15	NT2RP2002312	F-NT2RP2002312	2787	R-NT2RP2002312	7867
13	NT2RP2002316	F-NT2RP2002316	2788	R-NT2RP2002316	7868
	NT2RP2002325	F-NT2RP2002325	2789	R-NT2RP2002325	7869
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	NT2RP2002373	F-NT2RP2002373	2791		
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	NT2RP2002394	F-NT2RP2002394	2793	R-NT2RP2002394	7872
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	NT2RP2002426	F-NT2RP2002426	2795	R-NT2RP2002426	7874
25	NT2RP2002439	F-NT2RP2002439	2796	R-NT2RP2002439	7875
	NT2RP2002442	F-NT2RP2002442	2797		
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	NT2RP2002464	F-NT2RP2002464	2799	R-NT2RP2002464	7877
30	NT2RP2002475	F-NT2RP2002475	2800	R-NT2RP2002475	7878
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	NT2RP2002503	F-NT2RP2002503	2803	R-NT2RP2002503	7881
35	NT2RP2002504	F-NT2RP2002504	2804	R-NT2RP2002504	7882
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	NT2RP2002537	F-NT2RP2002537	2806	R-NT2RP2002537	7884
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40	NT2RP2002549	F-NT2RP2002549	2808	R-NT2RP2002549	7886
40	NT2RP2002591	F-NT2RP2002591	2809	R-NT2RP2002591	7887
	NT2RP2002595	F-NT2RP2002595	2810	R-NT2RP2002595	7888
	NT2RP2002606	F-NT2RP2002606	2811	R-NT2RP2002606	7889
	NT2RP2002609	F-NT2RP2002609	2812	R-NT2RP2002609	7890
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50	NT2RP2002701	F-NT2RP2002701	2817	R-NT2RP2002701	7895
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	NT2RP2002710	F-NT2RP2002710	2819	R-NT2RP2002710	7897
	NT2RP2002727	F-NT2RP2002727	2820	R-NT2RP2002727	7898
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	NT2RP2002740	F-NT2RP2002740	2822	R-NT2RP2002740	7900
	NT2RP2002741	F-NT2RP2002741	2823	R-NT2RP2002741	7901
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	NT2RP2002753	F-NT2RP2002753	2826	R-NT2RP2002753	7904
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	NT2RP2002778	F-NT2RP2002778	2828	R-NT2RP2002778	7906
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15	NT2RP2002880	F-NT2RP2002880	2833	R-NT2RP2002880	7911
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	NT2RP2002925	F-NT2RP2002925	2835	R-NT2RP2002925	7913
	NT2RP2002928	F-NT2RP2002928	2836	R-NT2RP2002928	7914
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20	NT2RP2002939	F-NT2RP2002939	2838		
	NT2RP2002954	F-NT2RP2002954	2839	R-NT2RP2002954	7916
	NT2RP2002959	F-NT2RP2002959	2840	R-NT2RP2002959	7917
	NT2RP2002979	F-NT2RP2002979	2841	R-NT2RP2002979	7918
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	NT2RP2003034	F-NT2RP2003034	2847	R-NT2RP2003034	7924
	NT2RP2003073	F-NT2RP2003073	2848	R-NT2RP2003073	7925
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	NT2RP2003117	F-NT2RP2003117	2851	R-NT2RP2003117	7928
	NT2RP2003121	F-NT2RP2003121	2852	R-NT2RP2003121	7929
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40	NT2RP2003129	F-NT2RP2003129	2854	R-NT2RP2003129	7931
40	NT2RP2003137	F-NT2RP2003137	2855	R-NT2RP2003137	7932
	NT2RP2003157	F-NT2RP2003157	2856		
	NT2RP2003158	F-NT2RP2003158	2857		
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45	NT2RP2003164	F-NT2RP2003164	2859	R-NT2RP2003164	7934
	NT2RP2003165	F-NT2RP2003165	2860	R-NT2RP2003165	7935
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50	NT2RP2003206	F-NT2RP2003206	2863	R-NT2RP2003206	7938
	NT2RP2003228	F-NT2RP2003228	2864		
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	NT2RP2003237	F-NT2RP2003237	2866	R-NT2RP2003237	7940
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	NT2RP2003293	F-NT2RP2003293	2873	R-NT2RP2003293	7947
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10	NT2RP2003297	F-NT2RP2003297	2875	R-NT2RP2003297	7949
	NT2RP2003307	F-NT2RP2003307	2876		
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	NT2RP3000092	F-NT2RP3000092	3187		
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	NT2RP3000267	F-NT2RP3000267	3202	R-NT2RP3000267	8259
	NT2RP3000299	F-NT2RP3000299,	3203	R-NT2RP3000299	8260
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	NT2RP3000348	F-NT2RP3000348	3209	R-NT2RP3000348	8266
25	NT2RP3000350	F-NT2RP3000350	3210	R-NT2RP30Q0350	8267
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		F-NT2RP3000562	3232	R-NT2RP3000562	8288
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	NT2RP3001432	F-NT2RP3001432	3333	R-NT2RP3001432	8385
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	NT2RP3001457	F-NT2RP3001457	3337	R-NT2RP3001457	8389
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30	NT2RP3001621	F-NT2RP3001621	3352	R-NT2RP3001621	8404
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	NT2RP3001634	F-NT2RP3001634	3354	R-NT2RP3001634	8406
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	NT2RP3001724	F-NT2RP3001724	3368	R-NT2RP3001724	8419
50	NT2RP3001727	F-NT2RP3001727	3369	n imonnocosaco	0.400
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5	NT2RP3001792	F-NT2RP3001792	3377	R-NT2RP3001792	8427
	NT2RP3001799	F-NT2RP3001799	3378	R-NT2RP3001799	8428
	NT2RP3001819	F-NT2RP3001819	3379	R-NT2RP3001819	8429
	NT2RP3001844	F-NT2RP3001844	3380	R-NT2RP3001844	8430
10	NT2RP3001854	F-NT2RP3001854	3381	R-NT2RP3001854	8431
	NT2RP3001855	F-NT2RP3001855	3382	R-NT2RP3001855	8432
	NT2RP3001857	F-NT2RP3001857	3383		
	NT2RP3001896	F-NT2RP3001896	3384	R-NT2RP3001896	8433
15	NT2RP3001898	F-NT2RP3001898	3385	R-NT2RP3001898	8434
,,	NT2RP3001915	F-NT2RP3001915	3386	R-NT2RP3001915	8435
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	NT2RP3001929	F-NT2RP3001929	3388	R-NT2RP3001929	8437
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	NT2RP3001969	F-NT2RP3001969	3393	R-NT2RP3001969	8442
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	NT2RP3002002	F-NT2RP3002002	3395	R-NT2RP3002002	8444
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	NT2RP3002033	F-NT2RP3002033	3399	R-NT2RP3002033	8448
	NT2RP3002045	F-NT2RP3002045	3400	R-NT2RP3002045	8449
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	NT2RP3002062	F-NT2RP3002062	3404	R-NT2RP3002062	8453
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	NT2RP3003833	F-NT2RP3003833	3555	R-NT2RP3003833	8593
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55	"ITHE DOOTED	1 WISH 2004419	5003	V 1417VL 2004412	0030

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13	NT2RP3004572	F-NT2RP3004572	3616	R-NT2RP3004572	8650
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50	NT2RP4000259	F-NT2RP4000259	3645	R-NT2RP4000259	8677
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	NT2RP4000290	F-NT2RP4000290	3647	R-nnnnnnnnnnn	8679
	NT2RP4000312	F-NT2RP4000312	3648	R-NT2RP4000312	8680
55	NT2RP4000321	F-NT2RP4000321	3649	R-NT2RP4000321	8681

	NT2RP4000323	F-NT2RP4000323	3650	R-NT2RP4000323	8682
	NT2RP4000355	F-NT2RP4000355	3651	R-NT2RP4000355	8683
	NT2RP4000360	F-NT2RP4000360	3652	R-NT2RP4000360	8684
5	NT2RP4000367	F-NT2RP4000367	3653	R-NT2RP4000367	8685
	NT2RP4000370	F-NT2RP4000370	3654	R-NT2RP4000370	8686
	NT2RP4000376	F-NT2RP4000376	3655	R-NT2RP4000376	8687
	NT2RP4000381	F-NT2RP4000381	3656	R-NT2RP4000381	8688
10	NT2RP4000398	F-NT2RP4000398	3657		
	NT2RP4000415	F-NT2RP4000415	3658	R-NT2RP4000415	8689
	NT2RP4000417	F-NT2RP4000417	3659	R-NT2RP4000417	8690
	NT2RP4000424	F-NT2RP4000424	3660	R-NT2RP4000424	8691
15	NT2RP4000448	F-NT2RP4000448	3661	R-NT2RP4000448	8692
,,	NT2RP4000449	F-NT2RP4000449	3662	R-NT2RP4000449	8693
	NT2RP4000455	F-NT2RP4000455,	3663	R-NT2RP4000455	8694
	NT2RP4000457	F-NT2RP4000457	3664	R-nnnnnnnnnnn	8695
	NT2RP4000480	F-NT2RP4000480	3665	R-NT2RP4000480	8696
20	NT2RP4000481	F-NT2RP4000481	3666	R-nnnnnnnnnnn	8697
	NT2RP4000498	F-NT2RP4000498	3667		
	NT2RP4000500	F-NT2RP4000500	3668	R-NT2RP4000500	8698
	NT2RP4000515	F-NT2RP4000515	3669	R-NT2RP4000515	8699
25	NT2RP4000517	F-NT2RP4000517	3670	R-NT2RP4000517	8700
	NT2RP4000518	F-NT2RP4000518	3671	R-NT2RP4000518	8701
	NT2RP4000519	F-NT2RP4000519	3672	R-NT2RP4000519	8702
	NT2RP4000524	F-NT2RP4000524	3673	R-NT2RP4000524	8703
30	NT2RP4000528	F-NT2RP4000528	3674	R-NT2RP4000528	8704
	NT2RP4000541	F-NT2RP4000541	3675	R-NT2RP4000541	8705
	NT2RP4000556	F-NT2RP4000556	3676	R-NT2RP4000556	8706
	NT2RP4000560	F-NT2RP4000560	3677		
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	NT2RP4000614	F-NT2RP4000614	3679	R-NT2RP4000614	8708
	NT2RP4000638	F-NT2RP4000638	3680	R-NT2RP4000638	8709
	NT2RP4000648	F-NT2RP4000648	3681	R-NT2RP4000648	8710
40	NT2RP4000657	F-NT2RP4000657	3682	R-NT2RP4000657	8711
,,,	NT2RP4000704	F-NT2RP4000704	3683	R-NT2RP4000704	8712
	NT2RP4000713	F-NT2RP4000713	3684		
	NT2RP4000724	F-NT2RP4000724	3685	R-NT2RP4000724	8713
	NT2RP4000728	F-NT2RP4000728	3686	R-NT2RP4000728	8714
45	NT2RP4000737	F-NT2RP4000737	3687		
	NT2RP4000739	F-NT2RP4000739	3688	R-NT2RP4000739	8715
	NT2RP4000781	F-NT2RP4000781	3689	R-NT2RP4000781	8716
	NT2RP4000787	F-NT2RP4000787	3690		
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	NT2RP4000833	F-NT2RP4000833	3692	R-NT2RP4000833	8718
	NT2RP4000837	F-NT2RP4000837	3693	R-NT2RP4000837	8719
	NT2RP4000839	F-NT2RP4000839	3694		
55	NT2RP4000855	F-NT2RP4000855	3695	R-NT2RP4000855	8720

		NT2RP4000865	F-NT2RP4000865	3696	R-NT2RP4000865	8721
		NT2RP4000878	F-NT2RP4000878	3697	R-NT2RP4000878	8722
		NT2RP4000879	F-NT2RP4000879	3698	R-NT2RP4000879	8723
	5	NT2RP4000907	F-NT2RP4000907	3699	R-nnnnnnnnnn	8724
		NT2RP4000915	F-NT2RP4000915	3700	R-nnnnnnnnnnn	8725
		NT2RP4000918	F-NT2RP4000918	3701		
		NT2RP4000925	F-NT2RP4000925	3702	R-NT2RP4000925	8726
	10	NT2RP4000927	F-NT2RP4000927	3703	R-nnnnnnnnnn	8727
		NT2RP4000928	F-NT2RP4000928	3704	R-NT2RP4000928	8728
		NT2RP4000929	F-NT2RP4000929	3705	R-NT2RP4000929	8729
		NT2RP4000955	F-NT2RP4000955	3706	R-NT2RP4000955	8730
	15	NT2RP4000973	F-NT2RP4000973	3707	R-NT2RP4000973	8731
		NT2RP4000975	F-NT2RP4000975	3708	R-NT2RP4000975	8732
		NT2RP4000979	F-NT2RP4000979,	3709	R-NT2RP4000979	8733
		NT2RP4000984	F-NT2RP4000984	3710	R-NT2RP4000984	8734
	20	NT2RP4000989	F-NT2RP4000989	3711	R-NT2RP4000989	8735
,	20	NT2RP4000996	F-NT2RP4000996	3712	R-NT2RP4000996	8736
		NT2RP4000997	F-NT2RP4000997	3713	R-NT2RP4000997	8737
		NT2RP4001004	F-NT2RP4001004	3714	R-NT2RP4001004	8738
		NT2RP4001006	F-NT2RP4001006	3715	R-NT2RP4001006	8739
•	25	NT2RP4001010	F-NT2RP4001010	3716	R-NT2RP4001010	8740
		NT2RP4001029	F-NT2RP4001029	3717	R-NT2RP4001029	8741
		NT2RP4001041	F-NT2RP4001041	3718	R-NT2RP4001041	8742
		NT2RP4001057	F-NT2RP4001057	3719	R-NT2RP4001057	8743
;	30	NT2RP4001064	F-NT2RP4001064	3720	R-NT2RP4001064	8744
		NT2RP4001078	F-NT2RP4001078	3721	R-NT2RP4001078	8745
		NT2RP4001079	F-NT2RP4001079	3722	R-NT2RP4001079	8746
		NT2RP4001080	F-NT2RP4001080	3723	R-NT2RP4001080	8747
;	35	NT2RP4001086	F-NT2RP4001086	3724	R-nnnnnnnnnnn	8748
		NT2RP4001095	F-NT2RP4001095	3725	R-NT2RP4001095	8749
		NT2RP4001100	F-NT2RP4001100	3726	R-NT2RP4001100	8750
		NT2RP4001117	F-NT2RP4001117	3727	R-NT2RP4001117	8751
4	10	NT2RP4001122	F-NT2RP4001122	3728	R-NT2RP4001122	8752
		NT2RP4001126	F-NT2RP4001126	3729	R-NT2RP4001126	8753
		NT2RP4001138	F-NT2RP4001138	3730	R-NT2RP4001138	8754
		NT2RP4001143	F-NT2RP4001143	3731	R-NT2RP4001143	8755
	<b>1</b> 5	NT2RP4001148	F-NT2RP4001148	3732	R-NT2RP4001148	8756
		NT2RP4001149	F-NT2RP4001149	3733	R-NT2RP4001149	8757
		NT2RP4001150	F-NT2RP4001150	3734	R-NT2RP4001150	8758
		NT2RP4001159	F-NT2RP4001159	3735	R-NT2RP4001159	8759
	-0	NT2RP4001174	F-NT2RP4001174	3736	R-NT2RP4001174	8760
•	50	NT2RP4001206	F-NT2RP4001206	3737	R-nnnnnnnnnn	8761
		NT2RP4001207	F-NT2RP4001207	3738	R-NT2RP4001207	8762
		NT2RP4001210	F-NT2RP4001210	3739	R-NT2RP4001210	8763
		NT2RP4001213	F-NT2RP4001213	3740	R-NT2RP4001213	8764
	55	NT2RP4001219	F-NT2RP4001219	3741	R-NT2RP4001219	8765

	NT2RP4001228	F-NT2RP4001228	3742	R-NT2RP4001228	8766
	NT2RP4001235	F-NT2RP4001235	3743	R-NT2RP4001235	8767
	NT2RP4001256	F-NT2RP4001256	3744	R-NT2RP4001256	8768
5	NT2RP4001260	F-NT2RP4001260	3745	R-NT2RP4001260	8769
	NT2RP4001274	F-NT2RP4001274	3746	R-NT2RP4001274	8770
	NT2RP4001276	F-NT2RP4001276	3747	R-nnnnnnnnnnnn	8771
	NT2RP4001313	F-NT2RP4001313	3748	R-NT2RP4001313	8772
10	NT2RP4001315	F-NT2RP4001315	3749	R-NT2RP4001315	8773
	NT2RP4001336	F-NT2RP4001336	3750	R NIZIG 4001010	00
	NT2RP4001339	F-NT2RP4001339	3751	R-NT2RP4001339	8774
	NT2RP4001343	F-NT2RP4001343	3752	N 11210 4001003	0
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15	NT2RP4001351	F-NT2RP4001351	3754	R-NT2RP4001351	8776
	NT2RP4001353	F-NT2RP4001353,	3755	R-NT2RP4001353	8777
	NT2RP4001372	F-NT2RP4001372			
	NT2RP4001373	F-NT2RP4001373	3756 3757	R-NT2RP4001372	8778
20	NT2RP4001375	F-NT2RP4001375		R-NT2RP4001373	8779
	NT2RP4001379	F-NT2RP4001379	3758	R-NT2RP4001375	8780
	NT2RP4001379	F-NT2RP4001379 F-NT2RP4001389	3759	R-NT2RP4001379	8781
	NT2RP4001389	F-NT2RP4001389	3760	R-NT2RP4001389	8782
25			3761	R-NT2RP4001407	8783
	NT2RP4001414	F-NT2RP4001414	3762	R-NT2RP4001414	8784
	NT2RP4001433	F-NT2RP4001433	3763	R-NT2RP4001433	8785
	NT2RP4001442	F-NT2RP4001442	3764	R-NT2RP4001442	8786
20	NT2RP4001447	F-NT2RP4001447	3765	R-NT2RP4001447	8787
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	NT2RP4001483	F-NT2RP4001483	3767	R-NT2RP4001483	8789
	NT2RP4001498	F-NT2RP4001498	3768	R-NT2RP4001498	8790
	NT2RP4001502	F-NT2RP4001502	3769	R-NT2RP4001502	8791
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	NT2RP4001524	F-NT2RP4001524	3771	R-NT2RP4001524	8793
	NT2RP4001529	F-NT2RP4001529	3772	R-NT2RP4001529	8794
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40	NT2RP4001551	F-NT2RP4001551	3774	R-nnnnnnnnnnn	8796
	NT2RP4001555	F-NT2RP4001555	3775	R-NT2RP4001555	8797
	NT2RP4001567	F-NT2RP4001567	3776	R-NT2RP4001567	8798
	NT2RP4001568	F-NT2RP4001568	3777	R-NT2RP4001568	8799
45	NT2RP4001571	F-NT2RP4001571	3778	R-NT2RP4001571	8800
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	NT2RP4001592	F-NT2RP4001592	3781	R-NT2RP4001592	8803
	NT2RP4001610	F-NT2RP4001610	3782	R-NT2RP4001610	8804
50	NT2RP4001614	F-NT2RP4001614	3783	R-NT2RP4001614	8805
	NT2RP4001634	F-NT2RP4001634	3784	R-NT2RP4001634	8806
	NT2RP4001638	F-NT2RP4001638	3785	R-NT2RP4001638	8807
	NT2RP4001644	F-NT2RP4001644	3786	R-NT2RP4001644	8808
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	NT2RP4001677	F-NT2RP4001677	3788	R-NT2RP4001677	8810
	NT2RP4001679	F-NT2RP4001679	3789		
	NT2RP4001696	F-NT2RP4001696	3790	R-NT2RP4001696	8811
5	NT2RP4001725	F-NT2RP4001725	3791	R-NT2RP4001725	8812
	NT2RP4001730	F-NT2RP4001730	3792	R-nnnnnnnnnnn	8813
	NT2RP4001739	F-NT2RP4001739	3793	R-NT2RP4001739	8814
	NT2RP4001753	F-NT2RP4001753	3794	R-NT2RP4001753	8815
10	NT2RP4001760	F-NT2RP4001760	3795	R-NT2RP4001760	8816
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	NT2RP4001803	F-NT2RP4001803	3797	R-NT2RP4001803	8818
	NT2RP4001822	F-NT2RP4001822	3798	R-NT2RP4001822	8819
15	NT2RP4001823	F-NT2RP4001823	3799	R-NT2RP4001823	8820
	NT2RP4001828	F-NT2RP4001828	3800	R-NT2RP4001828	8821
	NT2RP4001838	F-NT2RP4001838,	3801	R-NT2RP4001838	8822
	NT2RP4001841	F-NT2RP4001841	3802		
00	NT2RP4001849	F-NT2RP4001849	3803	R-NT2RP4001849	8823
20	NT2RP4001861	F-NT2RP4001861	3804		
	NT2RP4001889	F-NT2RP4001889	3805	R-NT2RP4001889	8824
	NT2RP4001893	F-NT2RP4001893	3806	R-NT2RP4001893	8825
	NT2RP4001896	F-NT2RP4001896	3807	R-NT2RP4001896	8826
25	NT2RP4001901	F-NT2RP4001901	3808	R-NT2RP4001901	8827
	NT2RP4001927	F-NT2RP4001927	3809	R-NT2RP4001927	8828
	NT2RP4001938	F-NT2RP4001938	3810	R-NT2RP4001938	8829
	NT2RP4001946	F-NT2RP4001946	3811	R-NT2RP4001946	8830
30	NT2RP4001950	F-NT2RP4001950	3812	R-NT2RP4001950	8831
	NT2RP4001953	F-NT2RP4001953	3813	R-NT2RP4001953	8832
	NT2RP4001966	F-NT2RP4001966	3814	R-NT2RP4001966	8833
	NT2RP4001975	F-NT2RP4001975	3815	R-NT2RP4001975	8834
35	NT2RP4002018	F-NT2RP4002018	3816	R-NT2RP4002018	8835
	NT2RP4002047	F-NT2RP4002047	3817	R-NT2RP4002047	8836
	NT2RP4002052	F-NT2RP4002052	3818	R-NT2RP4002052	8837
	NT2RP4002058	F-NT2RP4002058	3819	R-NT2RP4002058	8838
40	NT2RP4002071	F-NT2RP4002071	3820	R-NT2RP4002071	8839
	NT2RP4002075	F-NT2RP4002075	3821	R-NT2RP4002075	8840
	NT2RP4002078	F-NT2RP4002078	3822	R-NT2RP4002078	8841
	NT2RP4002081	F-NT2RP4002081	3823	R-nnnnnnnnnnn	8842
45	NT2RP4002083	F-NT2RP4002083	3824	R-NT2RP4002083	8843
	NT2RP4002408	F-NT2RP4002408	3825	R-NT2RP4002408	8844
	NT2RP4002791	F-NT2RP4002791	3826	R-NT2RP4002791	8845
	NT2RP4002888	F-NT2RP4002888	3827	R-NT2RP4002888	8846
50	NT2RP4002905	F-NT2RP4002905	3828	R-NT2RP4002905	8847
JU	NT2RP5003459	F-NT2RP5003459	3829		
	NT2RP5003461	F-NT2RP5003461	3830		
	NT2RP5003477	F-NT2RP5003477 F-NT2RP5003492	3831 3832		
	NT2RP5003492				
55	NT2RP5003500	F-NT2RP5003500	3833		

	NT2RP5003506	F-NT2RP5003506	3834		
	NT2RP5003512	F-NT2RP5003512	3835		
	NT2RP5003522	F-NT2RP5003522	3836		
5	NT2RP5003524	F-NT2RP5003524	3837		
	NT2RP5003534	F-NT2RP5003534	3838		
	OVARC1000001	F-0VARC1000001	3839	R-0VARC1000001	8848
	0VARC1000004	F-0VARC1000004	3840	R-0VARC1000004	8849
10	OVARC1000006	F-0VARC1000006	3841	R-0VARC1000006	8850
	OVARC1000013	F-0VARC1000013	3842	R-OVARC1000013	8851
	0VARC1000014	F-0VARC1000014	3843	R-OVARC1000014	8852
	0VARC1000017	F-0VARC1000017	3844	R-OVARC1000017	8853
15	OVARC1000035	F-0VARC1000035	3845	R-0VARC1000035	8854
	OVARC1000058	F-0VARC1000058	3846	R-0VARC1000058	8855
	OVARC1000060	F-0VARC1000060,	3847	R-0VARC1000060	8856
	0VARC1000068	F-0VARC1000068	3848	R-0VARC1000068	8857
20	OVARC1000071	F-0VARC1000071	3849	R-OVARC1000071	8858
	OVARC1000085	F-0VARC1000085	3850	R-0VARC1000085	8859
	OVARC1000087	F-0VARC1000087	3851	R-nnnnnnnnnnn	8860
	OVARC1000091	F-0VARC1000091	3852	R-0VARC1000091	8861
25	0VARC1000092	F-0VARC1000092	3853	R-0VARC1000092	8862
25	0VARC1000106	F-0VARC1000106	3854	R-0VARC1000106	8863
	0VARC1000109	F-0VARC1000109	3855		
	OVARC1000113	F-0VARC1000113	3856	R-0VARC1000113	8864
	OVARC1000114	F-0VARC1000114	3857	R-0VARC1000114	8865
30	0VARC1000133	F-0VARC1000133	3858	R-0VARC1000133	8866
	OVARC1000139 OVARC1000145	F-0VARC1000139	3859	D 0VADC1000145	8867
	0VARC1000145	F-0VARC1000145 F-0VARC1000148	3860 3861	R-0VARC1000145 R-0VARC1000148	8868
	0VARC1000148	F-0VARC1000148	3862	R-0VARC1000148	8869
35	0VARC1000151	F-0VARC1000151	3863	R-0VARC1000151	8870
	0VARC1000191	F-0VARC1000191	3864	R-0VARC1000191	8871
	0VARC1000191	F-0VARC1000198	3865	R-0VARC1000191	8872
	OVARC1000209	F-0VARC1000209	3866	R-0VARC1000209	8873
40	0VARC1000212	F-0VARC1000212	3867	R-0VARC1000212	8874
	0VARC1000240	F-0VARC1000240	3868	R-0VARC1000240	8875
	0VARC1000241	F-0VARC1000241	3869	R-0VARC1000241	8876
	OVARC1000288	F-0VARC1000288	3870	R-0VARC1000288	8877
45	0VARC1000302	F-0VARC1000302	3871	R-0VARC1000302	8878
	0VARC1000304	F-0VARC1000304	3872	R-0VARC1000304	8879
	0VARC1000309	F-0VARC1000309	3873	R-0VARC1000309	8880
	OVARC1000321	F-0VARC1000321	3874	R-0VARC1000321	8881
50	0VARC1000326	F-0VARC1000326	3875	R-0VARC1000326	8882
	0VARC1000335	F-0VARC1000335	3876	R-0VARC1000335	8883
	0VARC1000347	F-0VARC1000347	3877	R-0VARC1000347	8884
	0VARC1000384	F-0VARC1000384	3878	R-0VARC1000384	8885
55	0VARC1000408	F-0VARC1000408	3879	R-0VARC1000408	8886

	0VARC1000411	F-0VARC1000411	3880	R-OVARC1000411	8887
	OVARC1000414	F-0VARC1000414	3881	R-0VARC1000414	8888
	OVARC1000420	F-0VARC1000420	3882	R-0VARC1000420	8889
5	OVARC1000427	F-0VARC1000427	3883	R-OVARC1000427	8890
	0VARC1000431	F-0VARC1000431	3884	R-0VARC1000431	8891
	0VARC1000437	F-0VARC1000437	3885	R-0VARC1000437	8892
	OVARC1000440	F-0VARC1000440	3886	R-0VARC1000440	8893
10	0VARC1000442	F-0VARC1000442	3887	R-OVARC1000442	8894
	OVARC1000443	F-0VARC1000443	3888	R-0VARC1000443	8895
	OVARC1000461	F-0VARC1000461	3889	R-0VARC1000461	8896
	0VARC1000465	F-0VARC1000465	3890	R-0VARC1000465	8897
45	0VARC1000466	F-0VARC1000466	3891	R-0VARC1000466	8898
15	0VARC1000473	F-0VARC1000473	3892	R-OVARC1000473	8899
	0VARC1000479	F-0VARC1000479,	3893	R-OVARC1000479	8900
	0VARC1000486	F-0VARC1000486	3894	R-0VARC1000486	8901
	0VARC1000496	F-0VARC1000496	3895	R-0VARC1000496	8902
20	0VARC1000520	F-0VARC1000520	3896	R-OVARC1000520	8903
	0VARC1000526	F-0VARC1000526	3897	R-0VARC1000526	8904
	0VARC1000533	F-0VARC1000533	3898	R-0VARC1000533	8905
	OVARC1000543	F-0VARC1000543	3899	R-0VARC1000543	8906
25	0VARC1000556	F-0VARC1000556	3900	R-0VARC1000556	8907
	0VARC1000557	F-0VARC1000557	3901	R-0VARC1000557	8908
	0VARC1000564	F-0VARC1000564	3902	R-0VARC1000564	8909
	0VARC1000573	F-0VARC1000573	3903	R-OVARC1000573	8910
30	0VARC1000576	F-0VARC1000576	3904		
	0VARC1000578	F-0VARC1000578	3905	R-0VARC1000578	8911
	OVARC1000588	F-0VARC1000588	3906	R-0VARC1000588	8912
	0VARC1000605	F-0VARC1000605	3907	R-0VARC1000605	8913
35	0VARC1000622	F-0VARC1000622	3908	R-OVARC1000622	8914
	0VARC1000640	F-0VARC1000640	3909	R-0VARC1000640	8915
	OVARC1000649	F-0VARC1000649	3910		
	OVARC1000661	F-0VARC1000661	3911	R-0VARC1000661	8916
40	0VARC1000678	F-0VARC1000678	3912	R-0VARC1000678	8917
40	OVARC1000679	F-0VARC1000679	3913	R-nnnnnnnnnn	8918
	OVARC1000681	F-0VARC1000681	3914	R-OVARC1000681	8919
	0VARC1000682	F-0VARC1000682	3915		
	0VARC1000689	F-0VARC1000689	3916	R-0VARC1000689	8920
45	OVARC1000700	F-0VARC1000700	3917	R-OVARC1000700	8921
	OVARC1000703	F-0VARC1000703	3918	R-0VARC1000703	8922
	OVARC1000722	F-0VARC1000722	3919		
	OVARC1000730	F-0VARC1000730	3920	R-OVARC1000730	8923
50	OVARC1000746	F-0VARC1000746	3921	R-OVARC1000746	8924
	OVARC1000769	F-0VARC1000769	3922	R-OVARC1000769	8925
	0VARC1000771	F-OVARC1000771	3923	R-OVARC1000771	8926
	0VARC1000781	F-0VARC1000781	3924	R-OVARC1000781	8927
55	OVARC1000787	F-0VARC1000787	3925	R-OVARC1000787	8928

	OVARC1000800	F-0VARC1000800	3926	R-0VARC1000800	8929
	0VARC1000802	F-0VARC1000802	3927	R-0VARC1000802	8930
	0VARC1000834	F-0VARC1000834	3928	R-0VARC1000834	8931
5	OVARC1000846	F-0VARC1000846	3929	R-0VARC1000846	8932
	OVARC1000850	F-0VARC1000850	3930	R-0VARC1000850	8933
	0VARC1000862	F-0VARC1000862	3931	R-0VARC1000862	8934
	OVARC1000876	F-0VARC1000876	3932	R-0VARC1000876	8935
10	OVARC1000883	F-0VARC1000883	3933	R-0VARC1000883	8936
	OVARC1000885	F-0VARC1000885	3934	R-0VARC1000885	8937
	0VARC1000886	F-0VARC1000886	3935	R-0VARC1000886	8938
	OVARC1000890	F-0VARC1000890	3936		• • • • • • • • • • • • • • • • • • • •
15	0VARC1000891	F-0VARC1000891	3937	R-0VARC1000891	8939
73	OVARC1000897	F-0VARC1000897	3938	R-0VARC1000897	8940
	OVARC1000912	F-0VARC1000912,	3939	R-0VARC1000912	8941
	OVARC1000915	F-0VARC1000915	3940	R-0VARC1000915	8942
	OVARC1000924	F-0VARC1000924	3941	R-0VARC1000924	8943
20	0VARC1000936	F-0VARC1000936	3942	R-0VARC1000936	8944
	0VARC1000937	F-0VARC1000937	3943	R-0VARC1000937	8945
	OVARC1000945	F-0VARC1000945	3944	R-0VARC1000945	8946
	OVARC1000948	F-0VARC1000948	3945	R-0VARC1000948	8947
25	OVARC1000959	F-0VARC1000959	3946	R-0VARC1000959	8948
	0VARC1000960	F-0VARC1000960	3947	R-0VARC1000960	8949
	0VARC1000964	F-0VARC1000964	3948		
	OVARC1000971	F-0VARC1000971	3949	R-0VARC1000971	8950
30	OVARC1000984	F-0VARC1000984	3950	R-0VARC1000984	8951
	0VARC1000996	F-0VARC1000996	3951	R-0VARC1000996	8952
	OVARC1000999	F-0VARC1000999	3952	R-0VARC1000999	8953
	OVARC1001000	F-0VARC1001000	3953	R-0VARC1001000	8954
35	OVARC1001004	F-0VARC1001004	3954	R-0VARC1001004	8955
	OVARC1001010	F-0VARC1001010	3955	R-0VARC1001010	8956
	OVARC1001011	F-0VARC1001011	3956	R-0VARC1001011	8957
	OVARC1001032	F-0VARC1001032	3957	R-0VARC1001032	8958
40	OVARC1001034	F-0VARC1001034	3958	R-0VARC1001034	8959
40	OVARC1001038	F-0VARC1001038	3959	R-0VARC1001038	8960
	0VARC1001040	F-0VARC1001040	3960	R-0VARC1001040	8961
	OVARC1001044	F-0VARC1001044	3961	R-0VARC1001044	8962
	OVARC1001051	F-0VARC1001051	3962	R-0VARC1001051	8963
45	OVARC1001055	F-0VARC1001055	3963	R-0VARC1001055	8964
	OVARC1001062	F-0VARC1001062	3964	R-0VARC1001062	8965
	OVARC1001065	F-0VARC1001065	3965		
	OVARC1001068	F-0VARC1001068	3966	R-0VARC1001068	8966
50	OVARC1001072	F-0VARC1001072	3967	R-0VARC1001072	8967
	OVARC1001074	F-0VARC1001074	3968	R-0VARC1001074	8968
	OVARC1001085	F-0VARC1001085	3969	R-0VARC1001085	8969
	0VARC1001092	F-0VARC1001092	3970	R-0VARC1001092	8970
55	OVARC1001107	F-0VARC1001107	3971		

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	OVARC1001113	F-0VARC1001113	3972	R-0VARC1001113	8971
	OVARC1001117	F-0VARC1001117	3973	R-0VARC1001117	8972
	OVARC1001118	F-0VARC1001118	3974	R-0VARC1001118	8973
5	OVARC1001129	F-0VARC1001129	3975	R-OVARC1001129	8974
	OVARC1001154	F-0VARC1001154	3976		
	OVARC1001161	F-0VARC1001161	3977	R-0VARC1001161	8975
	OVARC1001162	F-0VARC1001162	3978	R-0VARC1001162	8976
10	OVARC1001167	F-0VARC1001167	3979	R-OVARC1001167	8977
	OVARC1001169	F-0VARC1001169	3980	R-OVARC1001169	8978
	OVARC1001170	F-0VARC1001170	3981	R-OVARC1001170	8979
	OVARC1001171	F-0VARC1001171	3982		
15	OVARC1001173	F-0VARC1001173	3983	R-OVARC1001173	8980
73	OVARC1001176	F-0VARC1001176	3984		
	OVARC1001180	F-0VARC1001180,	3985	R-0VARC1001180	8981
	OVARC1001188	F-0VARC1001188	3986	R-0VARC1001188	8982
	OVARC1001200	F-0VARC1001200	3987	R-0VARC1001200	8983
20	OVARC1001232	F-0VARC1001232	3988	R-OVARC1001232	8984
	OVARC1001240	F-0VARC1001240	3989	R-0VARC1001240	8985
	OVARC1001243	F-0VARC1001243	3990	R-0VARC1001243	8986
	OVARC1001244	F-0VARC1001244	3991		
25	OVARC1001261	F-0VARC1001261	3992	R-0VARC1001261	8987
	0VARC1001268	F-0VARC1001268	3993	R-0VARC1001268	8988
	OVARC1001270	F-0VARC1001270	3994	R-OVARC1001270	8989
	OVARC1001271	F-0VARC1001271	3995	R-OVARC1001271	8990
30	OVARC1001282	F-0VARC1001282	3996	R-OVARC1001282	8991
	0VARC1001296	F-0VARC1001296	3997	R-0VARC1001296	8992
	OVARC1001306	F-0VARC1001306	3998	R-nnnnnnnnnnn	8993
	OVARC1001329	F-0VARC1001329	3999	R-0VARC1001329	8994
25	OVARC1001330	F-0VARC1001330	4000	R-0VARC1001330	8995
35	OVARC1001339	F-0VARC1001339	4001	R-0VARC1001339	8996
	OVARC1001341	F-0VARC1001341	4002	R-0VARC1001341	8997
	OVARC1001342	F-0VARC1001342	4003	R-0VARC1001342	8998
	0VARC1001344	F-0VARC1001344	4004	R-0VARC1001344	8999
40	OVARC1001357	F-0VARC1001357	4005	R-0VARC1001357	9000
	OVARC1001360	F-0VARC1001360	4006	R-0VARC1001360	9001
	OVARC1001369	F-0VARC1001369	4007	R-0VARC1001369	9002
	OVARC1001372	F-0VARC1001372	4008	R-0VARC1001372	9003
45	OVARC1001376	F-0VARC1001376	4009	R-0VARC1001376	9004
	OVARC1001381	F-0VARC1001381	4010	R-0VARC1001381	9005
	OVARC1001391	F-0VARC1001391	4011	R-0VARC1001391	9006
	OVARC1001399	F-0VARC1001399	4012	R-nnnnnnnnnnn	9007
50	OVARC1001417	F-0VARC1001417	4013	R-0VARC1001417	9008
	OVARC1001419	F-0VARC1001419	4014	R-0VARC1001419	9009
	OVARC1001425	F-0VARC1001425	4015	R-0VARC1001425	9010
	OVARC1001436	F-0VARC1001436	4016	R-0VARC1001436	9011
<i>EE</i>	OVARC1001442	F-0VARC1001442	4017	R-0VARC1001442	9012
55					

	OVARC1001453	F-0VARC1001453	4018	R-0VARC1001453	- 9013
	0VARC1001476	F-0VARC1001476	4019	R-0VARC1001476	9014
	0VARC1001480	F-0VARC1001480	4020	R-0VARC1001480	9015
5	OVARC1001489	F-0VARC1001489	4021	R-0VARC1001489	9016
	0VARC1001496	F-0VARC1001496	4022	R-0VARC1001496	9017
	OVARC1001506	F-0VARC1001506	4023	R-0VARC1001506	9018
	OVARC1001525	F-0VARC1001525	4024	R-0VARC1001525	9019
10	OVARC1001542	F-0VARC1001542	4025	R-0VARC1001542	9020
	OVARC1001547	F-0VARC1001547	4026	R-0VARC1001547	9021
	OVARC1001555	F-0VARC1001555	4027		
	OVARC1001577	F-0VARC1001577	4028	R-OVARC1001577	9022
15	OVARC1001600	F-0VARC1001600	4029	R-0VARC1001600	9023
7.5	OVARC1001610	F-0VARC1001610	4030	R-0VARC1001610	9024
	OVARC1001611	F-0VARC1001611,	4031	R-0VARC1001611	9025
	OVARC1001615	F-0VARC1001615	4032	R-OVARC1001615	9026
	OVARC1001668	F-0VARC1001668	4033	R-0VARC1001668	9027
20	OVARC1001702	F-0VARC1001702	4034	R-0VARC1001702	9028
	OVARC1001703	F-0VARC1001703	4035	R-0VARC1001703	9029
	OVARC1001711	F-0VARC1001711	4036	R-0VARC1001711	9030
	OVARC1001713	F-0VARC1001713	4037		
25	OVARC1001726	F-0VARC1001726	4038	R-0VARC1001726	9031
	OVARC1001731	F-0VARC1001731	4039	R-0VARC1001731	9032
	OVARC1001745	F-0VARC1001745	4040	R-0VARC1001745	9033
	OVARC1001762	F-0VARC1001762	4041	R-nnnnnnnnnnn	9034
30	OVARC1001766	F-0VARC1001766	4042	R-0VARC1001766	9035
	OVARC1001767	F-0VARC1001767	4043	R-nnnnnnnnnnn	9036
	OVARC1001768	F-0VARC1001768	4044	R-0VARC1001768	9037
	OVARC1001791	F-0VARC1001791	4045	R-0VARC1001791	9038
35	OVARC1001795	F-0VARC1001795	4046	R-0VARC1001795	9039
33	OVARC1001802	F-0VARC1001802	4047	R-0VARC1001802	9040
	OVARC1001805	F-0VARC1001805	4048	R-0VARC1001805	9041
	OVARC1001809	F-0VARC1001809	4049		
	OVARC1001812	F-0VARC1001812	4050	R-0VARC1001812	9042
40	OVARC1001813	F-0VARC1001813	4051	R-0VARC1001813	9043
	OVARC1001820	F-0VARC1001820	4052	R-0VARC1001820	9044
	OVARC1001828	F-0VARC1001828	4053	R-0VARC1001828	9045
	OVARC1001846	F-0VARC1001846	4054	R-0VARC1001846	9046
45	OVARC1001861	F-0VARC1001861	4055	R-0VARC1001861	9047
	OVARC1001873	F-0VARC1001873	4056	R-0VARC1001873	9048
	OVARC1001879	F-0VARC1001879	4057	R-0VARC1001879	9049
	OVARC1001880	F-0VARC1001880	4058	R-0VARC1001880	9050
50	OVARC1001883	F-0VARC1001883	4059	R-0VARC1001883	9051
	OVARC1001900	F-0VARC1001900	4060	R-0VARC1001900	9052
	OVARC1001901	F-0VARC1001901	4061	R-0VARC1001901	9053
	OVARC1001911	F-0VARC1001911	4062	R-0VARC1001911	9054
55	OVARC1001916	F-0VARC1001916	4063	R-0VARC1001916	9055
33					

	OVARC1001928	F-0VARC1001928	4064	R-0VARC1001928	9056
	OVARC1001942	F-0VARC1001942	4065	R-0VARC1001942	9057
	OVARC1001943	F-0VARC1001943	4066	R-0VARC1001943	9058
5	OVARC1001949	F-0VARC1001949	4067	R-0VARC1001949	9059
	OVARC1001950	F-0VARC1001950	4068	R-0VARC1001950	9060
	OVARC1001987	F-0VARC1001987	4069	R-0VARC1001987	9061
	OVARC1001989	F-0VARC1001989	4070	R-0VARC1001989	9062
10	OVARC1002044	F-0VARC1002044	4071	R-0VARC1002044	9063
	OVARC1002050	F-0VARC1002050	4072	R-0VARC1002050	9064
	OVARC1002066	F-0VARC1002066	4073	R-0VARC1002066	9065
	OVARC1002082	F-0VARC1002082	4074	R-0VARC1002082	9066
	OVARC1002107	F-0VARC1002107	4075	R-0VARC1002107	9067
15	OVARC1002107	F-0VARC1002112	4076	R OTHEOTOGETOT	300.
	OVARC1002112	F-0VARC1002112	4077	R-0VARC1002127	9068
	OVARC1002127	F-0VARC1002138	4078	R-0VARC1002121	9069
	OVARC1002138	F-0VARC1002143	4079	R-0VARC1002143	9070
20	OVARC1002145	F-0VARC1002156	4080	R-0VARC1002145	9071
	OVARC1002158	F-0VARC1002158	4081	R-0VARC1002158	9072
	OVARC1002158	F-0VARC1002165	4082	R-0VARC1002165	9073
	0VARC1002183	F-0VARC1002182	4083	R-0VARC1002183	9074
25	PLACE1000004	F-PLACE1000004	4084	R-PLACE1000004	9075
	PLACE1000004	F-PLACE1000005	4085	R-PLACE1000005	9076
	PLACE1000007	F-PLACE1000007	4086	R-PLACE1000007	9077
	PLACE1000014	F-PLACE1000014	4087	R-PLACE1000014	9078
	PLACE1000014	F-PLACE1000014	4088	R-PLACE1000014	9079
30	PLACE1000031	F-PLACE1000031	4089	R-PLACE1000031	9080
	PLACE1000048	F-PLACE1000048	4090	R-PLACE1000040	9081
	PLACE1000048	F-PLACE1000048	4090	R-PLACE1000048	9082
	PLACE1000050	F-PLACE1000061	4091	R-PLACE1000050	9083
35	PLACE1000061	F-PLACE1000061	4092	R-PLACE1000061	9084
	PLACE1000078	F-PLACE1000078	4094	R-PLACE1000078	9085
	PLACE1000078	F-PLACE1000078	4095	R-PLACE1000078	9086
	PLACE1000081	F-PLACE1000081	4096	R-PLACE1000081	9087
40	PLACE1000133	F-PLACE1000034 F-PLACE1000133	4097	R-PLACE1000094	9088
	PLACE1000133	F-PLACE1000142	4098	R-PLACE1000142	9089
	PLACE1000142	F-PLACE1000142	4099	R-PLACE1000142	9090
	PLACE1000184	F-PLACE1000185	4100	R-PLACE1000184	9091
45	PLACE1000183	F-PLACE1000183	4100	R-PLACE1000183	9092
45	PLACE1000213	F-PLACE1000213	4101	R-PLACE1000213	9092
	PLACE1000214	F-PLACE1000214 F-PLACE1000236	4102	R-PLACE1000214	9094
	PLACE1000236	F-PLACE1000236	4103	R-PLACE1000236	9095
	PLACE1000240	F-PLACE1000248		R-PLACE1000240	9096
50	PLACE1000292 PLACE1000308	F-PLACE1000292 F-PLACE1000308	4105 4106	V_LTWOETOOOSAS	3030
	PLACE1000308	F-PLACE1000308	4106	R-PLACE1000332	9097
	PLACE1000332	F-PLACE1000332 F-PLACE1000347	4107	R-PLACE1000332	9098
	PLACE1000347 PLACE1000374	F-PLACE1000347	4109	R-PLACE1000347	9098
55	1 PUOP 1000 14	1 11400100014	4103	W 1 PUCETO00914	JUJJ

	PLACE1000380	F-PLACE1000380	4110	R-PLACE1000380	9100 -
	PLACE1000383	F-PLACE1000383	4111	R-PLACE1000383	9101
	PLACE1000401	F-PLACE1000401	4112	R-PLACE1000401	9102
5	PLACE1000406	F-PLACE1000406	4113	R-PLACE1000406	9103
	PLACE1000420	F-PLACE1000420	4114	R-PLACE1000420	9104
	PLACE1000421	F-PLACE1000421	4115	R-PLACE1000421	9105
	PLACE1000424	F-PLACE1000424	4116	R-PLACE1000424	9106
10	PLACE1000435	F-PLACE1000435	4117	R-PLACE1000435	9107
,,,	PLACE1000444	F-PLACE1000444	4118	R-PLACE1000444	9108
	PLACE1000453	F-PLACE1000453	4119	R-PLACE1000453	9109
	PLACE1000481	F-PLACE1000481	4120	R-PLACE1000481	9110
	PLACE1000492	F-PLACE1000492	4121	R-PLACE1000492	9111
15	PLACE1000540	F-PLACE1000432	4122	R-PLACE1000540	9112
	PLACE1000547	F-PLACE1000547,	4123	R-PLACE1000547	9113
	PLACE1000562	F-PLACE1000562	4123	R-PLACE1000547	9114
	PLACE1000564	F-PLACE1000564	4125	R-PLACE1000564	9115
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	PLACE1000588	F-PLACE1000588	4127		9117
	PLACE1000596	F-PLACE1000596	4128	R-nnnnnnnnnnn R-PLACE1000596	9118
	PLACE1000599	F-PLACE1000599	4129	R-PLACE1000599	9119
25	PLACE1000599	F-PLACE1000599			9119
20	PLACE1000611	F-PLACE1000610	4130	R-PLACE1000610	9120
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	PLACE1000656		4133	R-PLACE1000653	9122
30	PLACE100056	F-PLACE1000656	4134	R-PLACE1000656	9123
	PLACE1000708 PLACE1000712	F-PLACE1000706	4135	R-PLACE1000706	9124
		F-PLACE1000712	4136	R-PLACE1000712	9125
	PLACE1000716 PLACE1000748	F-PLACE1000716	4137	R-PLACE1000716	9126
35		F-PLACE1000748	4138	R-PLACE1000748	9127
	PLACE1000749	F-PLACE1000749	4139	R-PLACE1000749	9128
	PLACE1000755	F-PLACE1000755	4140	R-PLACE1000755	9129
	PLACE1000769	F-PLACE1000769	4141	R-PLACE1000769	9130
40	PLACE1000785	F-PLACE1000785	4142	R-PLACE1000785	9131
40	PLACE1000786	F-PLACE1000786	4143	R-PLACE1000786	9132
	PLACE1000793	F-PLACE1000793	4144	R-nnnnnnnnnnnn	9133
	PLACE1000798	F-PLACE1000798	4145	R-PLACE1000798	9134
	PLACE1000841	F-PLACE1000841	4146	R-PLACE1000841	9135
45	PLACE1000849	F-PLACE1000849	4147	R-nnnnnnnnnnn	9136
	PLACE1000856	F-PLACE1000856	4148	R-PLACE1000856	9137
	PLACE1000863	F-PLACE1000863	4149	R-PLACE1000863	9138
	PLACE1000909	F-PLACE1000909	4150	R-PLACE1000909	9139
50	PLACE1000931	F-PLACE1000931	4151	R-PLACE1000931	9140
	PLACE1000948	F-PLACE1000948	4152	R-PLACE1000948	9141
	PLACE1000972	F-PLACE1000972	4153	R-PLACE1000972	9142
	PLACE1000977	F-PLACE1000977	4154	R-PLACE1000977	9143
55	PLACE1000979	F-PLACE1000979	4155	R-PLACE1000979	9144
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	PLACE1001007	F-PLACE1001007	4158	R-PLACE1001007	9146
5	PLACE1001010	F-PLACE1001010	4159	R-PLACE1001010	9147
	PLACE1001015	F-PLACE1001015	4160	R-PLACE1001015	9148
	PLACE1001013	F-PLACE1001024	4161	R-PLACE1001024	9149
	PLACE1001024 PLACE1001036	F-PLACE1001024	4162	R-PLACE1001024	9150
		F-PLACE1001054		K-1 FVCF1001020	3130
10	PLACE1001054		4163	R-PLACE1001062	9151
	PLACE1001062	F-PLACE1001062 F-PLACE1001076	4164	R-PLACE1001002	9152
	PLACE1001076		4165	R-PLACE1001076	9153
	PLACE1001088	F-PLACE1001088	4166		
15	PLACE1001092	F-PLACE1001092	4167	R-PLACE1001092	9154
	PLACE1001104	F-PLACE1001104	4168	R-PLACE1001104	9155
	PLACE1001118	F-PLACE1001118,	4169	R-PLACE1001118	9156
	PLACE1001136	F-PLACE1001136	4170	R-PLACE1001136	9157
20	PLACE1001168	F-PLACE1001168	4171	R-PLACE1001168	9158
20	PLACE1001171	F-PLACE1001171	4172	R-PLACE1001171	9159
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55	PLACE1010947	F-PLACE1010947	4983	R-PLACE1010947	9943
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	THYRO100		YR01000895	5332	R-THYR01000895	10255
	THYRO100		YR01000916	5333	R-THYR01000916	10256
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5	Y79AA1001548	F-Y79AA1001548	5493	R-Y79AA1001548	10411
	Y79AA1001555	F-Y79AA1001555	5494	R-Y79AA1001555	10412
	Y79AA1001581	F-Y79AA1001581	5495		
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55	Y79AA1002258	F-Y79AA1002258	5534	R-Y79AA1002258	10450
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•	Y79AA1002361	F-Y79AA1002361	5539	R-Y79AA1002361	10455
	Y79AA1002399	F-Y79AA1002399	5540	R-Y79AA1002399	10456
	Y79AA1002407	F-Y79AA1002407	5541	R-Y79AA1002407	10457
10	Y79AA1002416	F-Y79AA1002416	5542	R-Y79AA1002416	10458
	Y79AA1002431	F-Y79AA1002431	5543	R-Y79AA1002431	10459
	Y79AA1002433	F-Y79AA1002433	5544	R-nnnnnnnnnnn	10460
	Y79AA1002472	F-Y79AA1002472	5545	R-Y79AA1002472	10461
15	Y79AA1002482	F-Y79AA1002482	5546	R-Y79AA1002482	10462
	Y79AA1002487	F-Y79AA1002487	5547	R-Y79AA1002487	10463

Table 2

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	name of	name of	SEQ ID	name of	SEQ ID
	clone	5'-end	of 5'-end	3'-end	of 3'-end
		sequence	sequence	sequence	sequence
10				•	
	HEMBA1000497	F-HEMBA100049	7 16111	R-HEMBA10004	97 16165
	HEMBA1001750	F-HEMBA100175	0 16112	R-HEMBA10017	50 16166
15	HEMBA1003854	F-HEMBA100385	4 16113	R-HEMBA10038	54 16167
	HEMBA1004193	F-HEMBA100419	3 16114	R-HEMBA10041	93 16168
	HEMBA1004860	F-HEMBA100486	0 16115	R-HEMBA10048	60 16169
	HEMBA1005572	F-HEMBA100557	2 16116	R-HEMBA10055	72 16170
20	HEMBA1006038	F-HEMBA100603	8 16117	R-HEMBA10060	38 16171
	HEMBA1006092	F-HEMBA100609	2 16118	R-HEMBA10060	92 16172
	HEMBA1006406	F-HEMBA100640	6 16119	R-HEMBA10064	06 16173
	HEMBA1006650	F-HEMBA100665	0 16120	R-HEMBA10066	50 16174
25	HEMBA1006812	F-HEMBA100681	2 16121	R-HEMBA10068	12 16175
	HEMBB1000672	F-HEMBB100067	2 16122	R-HEMBB10006	72 16176
	HEMBB1001197	F-HEMBB100119	7 16123	R-HEMBB10011	97 16177
	HEMBB1001871	F-HEMBB100187	1 16124	R-HEMBB10018	71 16178
30	MAMMA1001252	F-MAMMA100125	2 16125	R-MAMMA10012	52 16179
	MAMMA1002094	F-MAMMA100209	4 16126	R-MAMMA10020	94 16180
	NT2RM4000634	F-NT2RM400063	4 16127	R-NT2RM40006	34 16181
	NT2RM4000657	F-NT2RM400065	7 16128	R-NT2RM40006	57 16182
35	NT2RM4000783	F-NT2RM400078	3 16129	R-NT2RM40007	83 16183
	NT2RM4000857	F-NT2RM400085	7 16130	R-NT2RM40008	57 16184
	NT2RM4001178	F-NT2RM400117	8 16131	R-NT2RM40011	78 16185
40	NT2RM4002420	F-NT2RM400242	0 16132	R-NT2RM40024	20 16186
· •	NT2RP2000198	F-NT2RP200019	8 16133	R-NT2RP20001	98 16187
	NT2RP2000551	F-NT2RP200055	1 16134	R-NT2RP20005	51 16188
	NT2RP2000660	F-NT2RP200066	0 16135	R-NT2RP20006	60 16189

	NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
	NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
5	NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
-	NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
	NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
	NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
10	NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
	NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
	NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
	NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
15	NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
	NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
	NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
20	NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
20	NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
	NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
	NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206
25	NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
	NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
	NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
	NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
30	0VARC1000008	F-0VARC1000008	16157	R-OVARC1000008	16211
	0VARC1000724	F-0VARC1000724	16158	R-OVARC1000724	16212
	OVARC1000751	F-0VARC1000751	16159	R-0VARC1000751	16213
	OVARC1001029	F-0VARC1001029	16160	R-OVARC1001029	16214
35	PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
	PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
	PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
	PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218
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[0019] Furthermore, the present invention relates to the use of the above primers, as described below.

- (4) A polynucleotide which can be synthesized with the primer set of (2) or (3).
- (5) A polynucleotide comprising a coding region in the polynucleotide of (4).
- (6) A substantially pure protein encoded by polynucleotide of (4).
- (7) A partial peptide of the protein of (6).

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[0020] In addition, the present invention comprises a polynucleotide described below and a protein encoded by the polynucleotide.

- (8) An isolated polynucleotide selected from the group consisting of
  - (a) a polynucleotide comprising a coding region of the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
  - (b) a polynucleotide comprising a nucleotide sequence encoding a protein comprising the amino acid sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
  - (c) a polynucleotide comprising a nucleotide sequence encoding a protein comprising an amino acid sequence

selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, in which one or more amino acids are substituted, deleted, inserted, and/or added, wherein said protein is functionally equivalent to the protein comprising said amino acid sequence selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351;

- (d) a polynucleotide that hybridizes with a polynucleotide comprising a nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351, and that comprises a nucleotide sequence encoding a protein functionally equivalent to the protein encoded by the nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351;
- (e) a polynucleotide comprising a nucleotide sequence encoding a partial amino acid sequence of a protein encoded by the polynucleotide of (a) to (d);
- (f) a polynucleotide comprising a nucleotide sequence with at least 70% identity to the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351.
- (9). A substantially pure protein encoded by the polynucleotide of (8).
- (10) An antibody against the protein or peptide of any one of (6), (7), and (9).
- (11) A vector comprising the polynucleotide of (5) or (8).

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- (12) A transformant carrying the polynucleotide of (5) or (8), or the vector of (11).
- (13) A transformant expressively carrying the polynucleotide of (5) or (8), or the vector of (11).
- (14) A method for producing the protein or peptide of any one of (6), (7), and (9), comprising culturing the transformant of (13) and recovering the expression product.
- (15) An oligonucleotide comprising the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351 or the nucleotide sequence complementary to the complementary strand thereof, wherein said oligonucleotide comprises 15 nucleotides or more.
- (16) Use of the oligonucleotide of (15) as a primer for synthesizing a polynucleotide.
- (17) Use of the oligonucleotide of (15) as a probe for detecting a gene.
- (18) An antisense polynucleotide against the polynucleotide of (8), or the portion thereof.
- (19) A method for synthesizing a polynucleotide, the method comprising:
  - a) synthesizing a complementary strand using a cDNA library as a template, and using the primer set of (2) or (3), or the primer of (16); and
  - b) recovering the synthesized product.
- (20) The method of (19), wherein the cDNA library is obtainable by oligo-capping method.
- (21) The method of (19), wherein the complementary strand is obtainable by PCR.
- (22) A method for detecting the polynucleotide of (8), the method comprising:
  - a) incubating a target polynucleotide with the oligonucleotide of (15) under the conditions where hybridization occurs, and
  - b) detecting the hybridization of the target polynucleotide with the oligonucleotide of (15).
- (23) A database of polynucleotides and/or proteins, the database comprising information on at least one sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351 and/or the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, or a medium on which the database is stored.
- [0021] Any patents, patent applications, and publications cited herein are incorporated by reference.

### BRIEF DESCRIPTION OF THE DRAWINGS

- [0022] Figure 1 shows the restriction maps of vectors pME18SFL3 and pUC19FL3.
- [0023] Figure 2 shows the reproducibility of gene expression analysis. The respective intensities of gene expression observed in independent set of experiments are plotted in the vertical axis as well as in the horizontal axis.
- [0024] Figure 3 shows the detection limit in gene expression analysis. The intensity of expression is shown in the vertical axis and the concentration ( $\mu$ g/ml) of probe used is shown in the horizontal axis.

## 5 DETAILED DESCRIPTION OF THE INVENTION

[0025] Herein, "polynucleotide" is defined as a molecule in which multiple nucleotides are polymerized. There are no limitations in the number of the polymerized nucleotides. In case that the polymer contains relatively low number

of nucleotides, it is also described as an "oligonucleotide". The polynucleotide or the oligonucleotide of the present invention can be a natural or chemically synthesized product. Alternatively, it can be synthesized using a template DNA by an enzymatic reaction such as PCR.

[0026] All the cDNA provided by the invention are full-length cDNA. Herein, a "full-length cDNA" is defined as a cDNA which contains both ATG codon (the translation start site) and the stop codon. Accordingly, the untranslated regions, which are originally found in the upstream or downstream of the protein coding region in natural mRNA, may or may not be contained.

**[0027]** An "isolated polynucleotide" is a polynucleotide the structure of which is not identical to that of any naturally occurring nucleic acid or to that of any fragment of a naturally occurring genomic nucleic acid spanning more than three separate genes. The term therefore covers, for example,

- (a) a DNA which has the sequence of part of a naturally occurring genomic DNA molecule but is not flanked by both of the coding sequences that flank that part of the molecule in the genome of the organism in which it naturally occurs;
- (b) a nucleic acid incorporated into a vector or into the genomic DNA of a prokaryote or eukaryote in a manner such that the resulting molecule is not identical to any naturally occurring vector or genomic DNA:

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- (c) a separate molecule such as a cDNA, a genomic fragment, a fragment produced by polymerase chain reaction (PCR), or a restriction fragment; and
- (d) a recombinant nucleotide sequence that is part of a hybrid gene, i.e., a gene encoding a fusion protein. Specifically excluded from this definition are nucleic acids present in mixtures of different (i) DNA molecules, (ii) transfected cells, or (iii) cell clones: e.g., as these occur in a DNA library such as a cDNA or genomic DNA library.

[0028] The term "substantially pure" as used herein in reference to a given polypeptide means that the protein or polypeptide is substantially free from other biological macromolecules. The substantially pure protein or polypeptide is at least 75% (e.g., at least 80, 85, 95, or 99%) pure by dry weight. Purity can be measured by any appropriate standard method, for example, by column chromatography, polyacrylamide gel electrophoresis, or HPLC analysis.

[0029] All the clones (5602 clones) of the present invention are novel and encode the full-length proteins. All the clones were prepared by oligo capping method, which can achieve cDNA cloning with high fullness ratio. The cDNA clones were selected by using ATGprl score as an index of the fullness ratio at the 5'-end, based on the sequence features of the 5'-end sequences. Selection was further carried out by searching GenBank database for EST sequences homologous to 5'-end sequence of each clone by BLAST [S.F. Altschul, W. Gish, W. Miller, E.W. Myers & D.L. Lipman J. Mol. Biol., 215:403-410 (1990); W. Gish, & D.J. States, Nature Genet., 3:266-272 (1993)] and by considering the number of matching (identical) EST sequences or the number of continuous amino acids in the 5'-end sequence initiated from the initiation codon.

[0030] Moreover, the clones were turn out to be not identical to any of the known human mRNA (namely novel) by homology search using the 5'-end sequence.

[0031] The primers of the present invention, which are used for synthesizing full-length cDNA, are selected from the group comprising SEQ ID NO: 1-5547 (5'-primer), or SEQ ID NO: 5548-10463 (3'-primer). Further, the primers of the present invention, which are used for synthesizing full-length cDNA, are selected from SEQ ID NO: 16111-16164 (5'-primer), or SEQ ID NO: 16165-16218 (3'-primer). Some of the nucleotides include a known EST as its part. However, the primers of the present invention are novel in terms that the primers enable to synthesize full-length cDNA. Because the known ESTs lack important information on what part of cDNA the ESTs correspond to, it is impossible to design primers on the basis of the ESTs.

[0032] All the full-length cDNA of the present invention can be synthesized using a primer set comprising the nucleotide sequences selected from both the 5'-and 3'-end sequences, or a set comprising a primer based on the 5'-end sequence and an oligo-dT primer, by a method such as PCR (Current protocols in Molecular Biology (1987) Ausubel et al. edit, John Wiley & Sons, Section 6.1-6.4).

[0033] Specifically, PCR can be performed using an oligonucleotide that has 15 nucleotides longer, and specifically hybridizes with the complementary strand of the polynucleotide that contains the nucleotide sequence selected from the 5'-end sequences shown in Table 1 and 2 (SEQ ID NO: 1-5547, or SEQ ID NO: 16111-16164), and an oligo-dT primer as a 5'-, and 3'-primer, respectively. The length of the primers is usually 15-100 bp, and favorably between 15-35 bp. In case of LA PCR, which is described below, the primer length of 25-35 bp may provide a good result.

[0034] A method to design a primer that enables a specific amplification based on the given nucleotide sequence is known to those skilled in the art (Current Protocols in Molecular Biology, Ausubel et al. edit, (1987) John Wiley & Sons, Section 6.1-6.4). In designing a primer based on the 5'-end sequence, the primer is designed so as that, in principle, the amplification products will include the translation start site. Accordingly, in case that a given 5'-end nucleotide sequence is the 5'- untranslated region (5'UTR), any part of the sequence can be used as a 5'-primer as far as the specificity toward the target cDNA is insured. The translation start site can be predicted using a known method such

as the ATGpr as described below.

[0035] When synthesizing a polynucleotide, the target nucleotide sequence to be amplified can extend to several thousand bp in some cDNA. However, it is possible to amplify such a long nucleotides by using such as LA PCR (Long and Accurate PCR). It is advantageous to use LA PCR when synthesizing long DNA. In LA PCR, in which a special DNA polymerase having 3'→5' exonuclease activity is used, misincorporated nucleotides can be removed. Accordingly, accurate synthesis of the complementary strand can be achieved even with a long nucleotide sequence. By using LA PCR, it is reported that amplification of a nucleotide with 20 kb longer can be achieved under desirable condition (Takeshi Hayashi (1996) Jikken-Igaku Bessatsu, "Advanced Technologies in PCR" Youdo-sha).

[0036] A template DNA for synthesizing the cDNA of the present invention can be obtained by using cDNA libraries that are prepared by various methods. The full-length cDNA clones obtained here are those with high fullness ratio, which were obtained using a combination of (1) a method to prepare a full-length-enriched cDNA library using the oligo-capping method, and (2) an estimation system for fullness using the 5'-end sequence (selection based on the estimation by the ATGpr after removing clones that are not-full-length compared to the ESTs). However, it is possible to easily obtain a full-length cDNA by using the primers that are provided by the present invention, not by the above described specialized method.

The problem with the cDNA libraries prepared by the known methods or commercially available is that mRNA contained in the libraries has very low fullness ratio. Thus, it is difficult to screen full-length cDNA clone directly from the library using ordinary cloning methods. The present invention has revealed a primer that is capable of synthesizing a full-length cDNA. If provided with primers, it is possible to synthesize a target full-length cDNA by using enzymatic reactions such as PCR. In particular, a full-length-enriched cDNA library, synthesized by methods such as oligo-capping, is desirable to synthesize a full-length cDNA with more reliability.

[0037] The 5'-end sequence of the full-length cDNA clones of the invention can be used to isolate the regulatory element of transcription including the promoter on the genome. By the spring of the year 2000, a rough draft of the human genome (analysis of human genomic sequence with lower accuracy), which covers 90% of the genome, is planned to be accomplished, and by the year 2003, analysis of the entire human genomic sequence is going to be finished. However, it is hard to analyze with software the transcription start sites on the human genome, in which long introns exist. By contrast, it is easy to specify the transcription start site on the genomic sequence using the 5'-end sequence of the full-length cDNA clone, thus it is easy to obtain the genomic region involved in transcription regulation, which includes the promoter that is contained in the upstream of the transcription start site.

[0038] The full-length cDNAs cloned in the present invention are classified into 13 groups, based on the data such as ATGpr1 score, by which the fullness ratio can be evaluated. Specifically, the 13 groups consist of; the below-mentioned groups (1)-(3), containing 3690 clones (Table 9), and the group (12), containing 3 clones, wherein ATGpr1 (score defined in the ATGpr program) is higher than 0.3; and the below-mentioned groups (4)-(11), containing 1857 clones (Table 10), and the group (13), containing 52 clones, wherein, although ATGpr1 is 0.3 or less, the clones are judged to be full-length from various viewpoints. Names of the clones belonging to the groups (1)-(13) are as indicated in Examples or below.

### (1) 1516 clones

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Among the 3690 clones that have the maximal ATGpr1 score higher than 0.3, 1516 clones are novel full-length clones, in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

#### (2)377 clones

Among the 3690, 377 clones are novel full-length clones, in which the number of human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

#### (3) 1797 clones

Among the 3690, 1797 clones are novel full-length clones, in which the number of human EST having identical sequence at the 5'-end is not more than 20 (except the clones described above).

### (4) 453 clones

Among the 1857 clones in which the maximal ATGpr1 score is 0.3 or less, the following 453 clones are estimated to be novel full-length clones since the clones have the maximal score 0.3 or more in the ATGpr2, and at least either of the sequences of their 5'- and 3'-ends, or both are not identical to those of any human EST. The ATGpr2 score is determined by using the ATGpr program with neglecting the information of the frequency of the six nucleotides contained within the sequence between the ATG codon and the stop codon (the maximal length is 300 nucleotides from the ATG codon) (Salamo A.A., Nishikawa T., and Swindells M.B. (1998) Bioinformatics, 14: 384-390; http://www.hri.co.jp/atgpr/). The ATGpr program for calculating the ATGpr2 score is described as the

ATGpr2 program in the followings.

#### (5) 24 clones

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Among the 1857 clones, 24 clones are estimated to be full-length since their maximal ATGpr2 scores are higher than 0.3, and also novel, though they have low scores in ATGpr1 program, in which the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

### (6) 65 clones

Among the 1857 clones, 65 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

#### (7) 32 clones

Among the 1857 clones, 32 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

#### (8) 36 clones

Among the 1857 clones, 36 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

### (9) 81 clones

Among the 1857 clones, 81 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if the number of the human EST having identical sequence at the 5'-end is not more than 20 (other than the clones in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST).

### (10) 938 clones

Among the 1857 clones, 938 clones are estimated to be full-length according to the fullness ratio shown in Table 4, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 5'-end is not identical to those of any human EST.

### (11) 228 clones

Among the 1857 clones, 228 clones are estimated to be full-length according to the fullness ratio shown in Table 7, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 3'-end is not identical to those of any human EST.

#### (12) 3 clones

Three clones, HEMBA1006812, HEMBB1001871, and NT2RP3001282, whose maximal ATGpr1 values are higher than 0.3, are full-length and novel clones whose 5'-end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 100 amino acids or more.

#### (13) 52 clones

The following 52 clones, which have maximal ATGpr1 values of 0.3 or less, are full-length with the fullness ratios shown in Table 4 although the fullness ratios are low:

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HEMBA1000497, HEMBA1001750, HEMBA1003854,
                                           HEMBA1004193, HEMBA1004860,
                                                                         HEMBA1005572.
HEMBA1006038, HEMBA1006092,
                             HEMBA1006406, HEMBA1006650,
                                                                          HEMBB1001197,
                                                           HEMBB1000672,
MAMMA1001252, MAMMA1002094, NT2RM4000634, NT2RM4000657,
                                                                          NT2RM4000857,
                                                           NT2RM4000783,
NT2RM4001178, NT2RM4002420,
                             NT2RP2000198,
                                            NT2RP2000551,
                                                           NT2RP2000660,
                                                                          NT2RP2001214,
NT2RP2001460, NT2RP2001756,
                             NT2RP2002056.
                                            NT2RP2002677,
                                                           NT2RP2002755.
                                                                          NT2RP2002843,
NT2RP2003101,
              NT2RP2003799,
                             NT2RP2004095,
                                            NT2RP2004732,
                                                           NT2RP2004920,
                                                                          NT2RP2005454.
NT2RP2005776, NT2RP2005806,
                             NT2RP2005882,
                                                           NT2RP3002099,
                                                                          NT2RP3003155,
                                            NT2RP3001723,
NT2RP3004028, OVARC1000008,
                             OVARC1000724, OVARC1000751,
                                                           OVARC1001029,
                                                                          PLACE1000814,
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PLACE1003030, PLACE1005549, PLACE1007218, NT2RP4002298.

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Moreover, the clones are novel clones whose 5' -end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 50 amino acids or more. Among them, the following 20 clones is predicted to contain a coding region with 100 amino acids or more and should encode proteins:

HEMBA1000497, HEMBA1003854, HEMBA1004193, NT2RM4000657, NT2RM4001178, NT2RP2001756, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2004095, NT2RP2004920, NT2RP2005806, NT2RP3002099, NT2RP3003155, OVARC1000724, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218.

[0039] The protein encoded by the polynucleotide of the invention can be prepared as a recombinant protein or as a natural protein. For example, the recombinant protein can be prepared by inserting the polynucleotide encoding the protein of the invention into a vector, introducing the vector into an appropriate host cell and purifying the protein expressed within the transformed host cell, as described below. In contrast, the natural protein can be prepared, for example, by utilizing an affinity column to which an antibody against the protein of the invention (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 16.1-16.19) is attached. The antibody used for affinity purification may be either a polyclonal antibody, or a monoclonal antibody. Alternatively, in vitro translation (See, for example, "On the fidelity of mRNA translation in the nuclease-treated rabbit reticulocyte lysate system." Dasso M.C., and Jackson R.J. (1989) Nucleic Acids Res. 17: 3129-3144) may be used for preparing the protein of the invention. [0040] Proteins functionally equivalent to the proteins of the present invention can be prepared based on the activities, which were clarified in the above-mentioned manner, of the proteins of the present invention. Using the biological activity possessed by the protein of the invention as an index, it is possible to verify whether or not a particular protein is functionally equivalent to the protein of the invention by examining whether or not the protein has said activity.

[0041] Proteins functionally equivalent to the proteins of the present invention can be prepared by those skilled in the art, for example, by using a method for introducing mutations into an amino acid sequence of a protein (for example, site-directed mutagenesis (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 8.1-8.5). Besides, such proteins can be generated by spontaneous mutations. The present invention comprises the proteins having one or more amino acids substitutions, deletions, insertions and/or additions in the amino acid sequences of the proteins of the present invention (Tables 350 and 351), as far as the proteins have the equivalent functions to those of the proteins identified in the present Examples described later.

[0042] There are no limitations in the number and sites of amino acid mutations, as far as the proteins maintain the functions thereof. The number of mutations is typically 30% or less, or 20% or less, or 10% or less, preferably within 5% or less, or 3% or less of the total amino acids, more preferably within 2% or less or 1 % or less of the total amino acids. From the viewpoint of maintaining the protein function, it is preferable that a substituted amino has a similar property to that of the original amino acid. For example, Ala, Val, Leu, Ile, Pro, Met, Phe and Trp are assumed to have similar properties to one another because they are all classified into a group of non-polar amino acids. Similarly, substitution can be performed among non-charged amino acid such as Gly, Ser, Thr, Cys, Tyr, Asn, and Gln, acidic amino acids such as Asp and Glu, and basic amino acids such as Lys, Arg, and His.

[0043] In addition, proteins functionally equivalent to the proteins of the present invention can be isolated by using techniques of hybridization or gene amplification known to those skilled in the art. Specifically, using the hybridization technique (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)), those skilled in the art can usually isolate a DNA highly homologous to the DNA encoding the protein identified in the present Example based on the identified nucleotide sequence (Tables 350 and 351) or a portion thereof and obtain the functionally equivalent protein from the isolated DNA. The present invention include proteins encoded by the DNAs hybridizing with the DNAs encoding the proteins identified in the present Example, as far as the proteins are functionally equivalent to the proteins identified in the present Example. Organisms from which the functionally equivalent proteins are isolated are illustrated by vertebrates such as human, mouse, rat, rabbit, pig and bovine, but are not limited to these animals.

[0044] Washing conditions of hybridization for the isolation of DNAs encoding the functionally equivalent proteins are usually " $1 \times SSC$ , 0.1% SDS, 37°C"; more stringent conditions are " $0.5 \times SSC$ , 0.1% SDS, 42°C"; and still more stringent conditions are " $0.1 \times SSC$ , 0.1% SDS, 65°C". Alternatively, the following conditions can be given as hybridization conditions of the present invention. Namely, conditions in which the hybridization is done at " $6 \times SSC$ , 40% Formamide, 25°C", and the washing at " $1 \times SSC$ , 55°C" can be given. More preferable conditions are those in which the hybridization is done at " $6 \times SSC$ , 40% Formamide, 37°C", and the washing at " $0.2 \times SSC$ , 55°C". Even more preferable are those in which the hybridization is done at " $6 \times SSC$ , 50% Formamide, 37°C", and the washing at " $0.1 \times SSC$ , 62°C". The more stringent the conditions of hybridization are, the more frequently the DNAs highly homologous to the probe sequence are isolated. Therefore, it is preferable to conduct hybridization under stringent conditions. Examples of stringent conditions in the present invention are, washing conditions of " $0.5 \times SSC$ , 0.1% SDS, 0.1% SDS

Those skilled in the art can select the hybridization conditions with similar stringency to those mentioned above by properly combining the above-mentioned or other factors (for example, probe concentration, probe length and duration of hybridization reaction) that determines the stringency of hybridization.

[0045] The amino acid sequences of proteins isolated by using the hybridization techniques usually exhibit high homology to those of the proteins of the present invention, which are shown in Tables 350 and 351. The present invention encompasses a polynucleotide comprising a nucleotide sequence that has a high identity to the nucleotide sequence of claim 8 (a).

Furthermore, the present invention encompasses a peptide, or protein comprising an amino acid sequence that has a high identity to the amino acid sequence encoded by the polynucleotide of claim 8 (b). The term "high identity" indicates sequence identity of at least 40% or more;

preferably 60% or more; and more preferably 70% or more. Alternatively, more preferable is identity of 90% or more, or 93% or more, or 95% or more, furthermore, 97% or more, or 99% or more. The identity can be determined by using the BLAST search algorithm.

[0046] With the gene amplification technique (PCR) (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)) using primers designed based on the nucleotide sequence (Tables 350 and 351) or a portion thereof identified in the present Example, it is possible to isolate a DNA fragment highly homologous to the polynucleotide sequence or a portion thereof and to obtain functionally equivalent protein to a particular protein identified in the present Example based on the isolated DNA fragment.

[0047] The "percent identity" of two amino acid sequences or of two nucleic acids is determined using the algorithm of Karlin and Altschul (Proc. Natl. Acad. Sei. USA 87:2264-2268, 1990), modified as in Karlin and Altschul (Proc. Natl. Acad. Sei. USA 90:5873-5877, 1993). Such an algorithm is incorporated into the BLASTN and BLASTX programs of Altschul et al. (J. Mol. Biol.215:403-410, 1990). BLAST nucleotide searches are performed with the BLASTN program, score = 100, wordlength = 12. BLAST protein searches are performed with the BLASTX program, score = 50, wordlength = 3. When gaps exist between two sequences, Gapped BLAST is utilized as described in Altschul et al. (Nucleic Acids Res.25:3389-3402,1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., BLASTX and BLASTN) are used. See http://www.ncbi.nlm.nih.gov.

[0048] The present invention also includes a partial peptide of the proteins of the invention. The partial peptide comprises a protein generated as a result that a signal peptide has been removed from a secretory protein. If the protein of the present invention has an activity as a receptor or a ligand, the partial peptide may function as a competitive inhibitor of the protein and may bind to the receptor (or ligand). In addition, the present invention comprises an antigen peptide for raising antibodies. For the peptides to be specific for the protein of the invention, the peptides comprise at least 7 amino acids, preferably 8 amino acids or more, more preferably 9 amino acids or more, and even more preferably 10 amino acids or more. The peptide can be used for preparing antibodies against the protein of the invention, or competitive inhibitors of them, and also screening for a receptor that binds to the protein of the invention. The partial peptides of the invention can be produced, for example, by genetic engineering methods, known methods for synthesizing peptides, or digesting the protein of the invention with an appropriate peptidase.

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[0049] The present invention also relates to a vector into which the DNA of the invention is inserted. The vector of the invention is not limited as long as it contains the inserted DNA stably. For example, if *E. coli* is used as a host, vectors such as pBluescript vector (Stratagene) are preferable as a cloning vector. To produce the protein of the invention, expression vectors are especially useful. Any expression vector can be used as far as it is capable of expressing the protein *in vitro*, in *E. coli*, in cultured cells, or *in vivo*. For example, pBEST vector (Promega) is preferable for *in vitro* expression, pET vector (Invitrogen) for *E. coli*, pME18S-FL3 vector (GenBank Accession No. AB009864) for cultured cells, and pME18S vector (Mol. Cell. Biol. (1988) 8: 466-472) for *in vivo* expression. To insert the DNA of the invention, ligation utilizing restriction sites can be performed according to the standard method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

[0050] The present invention also relates to a transformant carrying the vector of the invention. Any cell can be used as a host into which the vector of the invention is inserted, and various kinds of host cells can be used depending on the purposes. For strong expression of the protein in eukaryotic cells, COS cells or CHO cells can be used, for example. [0051] Introduction of the vector into host cells can be performed, for example, by calcium phosphate precipitation method, electroporation method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 9.1-9.9), lipofectamine method (GIBCO-BRL), or microinjection method, etc.

[0052] The primer of the present invention can be used for synthesizing full-length cDNA, and also for the detection and/or diagnosis of the abnormality of the protein of the invention encoded by the full-length cDNA. For example, by utilizing polymerase chain reaction (genomic DNA-PCR, or RT-PCR) using the primer of the invention, DNA encoding the protein of the invention can be amplified. It is also possible to obtain the regulatory region of expression in the 5'-upstream by using PCR or hybridization since the transcription start site within the genomic sequence can be easily specified based on the 5'-end sequence of the full-length cDNA. The obtained genomic region can be used for detection and/or diagnosis of the abnormality of the sequence by RFLP analysis, SSCP, or direct sequencing.

[0053] Furthermore, the "polynucleotide having a length of at least 15 nucleotides, comprising a nucleotide sequence that is complementary to a polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs in Tables 350 and 351, or its complementary strand" includes an antisense polynucleotide for suppressing the expression of the protein of the invention. To exert the antisense effect, the antisense polynucleotide has a length of at least 15 bp or more, for example, 50 bp or more, preferably 100 bp or more, and more preferably 500 bp or more, and has a length of usually 3000 bp or less and preferably 2000 bp or less. The antisense DNA can be used in the gene therapy of the diseases that are caused by the abnormality of the protein of the invention (abnormal function or abnormal expression). Said antisense DNA can be prepared, for example, by the phosphorothioate method ("Physicochemical properties of phosphorothioate oligodeoxynucleotides." Stein (1988) Nucleic Acids Res. 16: 3209-3221) based on the nucleotide sequence of the DNA encoding the protein (for example, the DNA set forth in any one of SEQ ID NOs in Tables 350 and 351).

[0054] The polynucleotide or antisense DNA of the present invention can be used in gene therapy, for example, by administrating it into a patient by the in vivo or ex vivo method with virus vectors such as retrovirus vectors, adenovirus vectors, and adeno-associated virus vectors, or non-virus vectors such as liposome.

[0055] The present invention also relates to antibodies that bind to the protein of the invention.

There are no limitations in the form of the antibodies of the invention. They include polyclonal antibodies, monoclonal antibodies, or their portions that can bind to the protein of the invention. They also include antibodies of all classes. Furthermore, special antibodies such as humanized antibodies are also included.

[0056] The polyclonal antibody of the invention can be obtained according to the standard method by synthesizing an oligopeptide corresponding to the amino acid sequence and immunizing rabbits with the peptides (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.12-11.13). The monoclonal antibody of the invention can be obtained according to the standard method by purifying the protein expressed in *E. coli*, immunizing mice with the protein, and producing a hybridoma cell by fusing the spleen cells and myeloma cells (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

[0057] The antibody binding to the protein of the present invention can be used for purification of the protein of the invention, and also for detection and/or diagnosis of the abnormalities of the expression and structure of the protein. Specifically, proteins can be extracted, for example, from tissues, blood, or cells, and the protein of the invention is detected by Western blotting, immunoprecipitation, or ELISA, etc. for the above purpose.

[0058] Furthermore, the antibody binding to the protein of the present invention can be utilized for treating the diseases that associates with the protein of the invention. If the antibodies are used for treating patients, human antibodies or humanized antibodies are preferable in terms of their low antigenicity. The human antibodies can be prepared by immunizing a mouse whose immune system is replaced with that of human ("Functional transplant of megabase human immunoglobulin loci recapitulates human antibody response in mice" Mendez M.J. et al. (1997) Nat. Genet. 15: 146-156). The humanized antibodies can be prepared by recombination of the hypervariable region of a monoclonal antibody (Methods in Enzymology (1991) 203: 99-121).

[0059] The cDNA of the present invention encodes the amino acid sequence of a protein which is predicted to have the function(s) described below based on the homology search of the GenBank and SwissProt. Specifically, for instance, as shown in EXAMPLES, searching a known gene or protein that is homologous to the partial sequence of the full-length cDNA of the invention (5602 clone) and referring the function of the gene and of the protein encoded by the gene make it possible to predict the function of the protein encoded by the cDNA of the invention. In this way, each of 1437 clones out of the 5602 full-length cDNA clones of the invention was predicted to encode a protein that was classified into one or more of the following categories.

Secretory or membrane protein (261 clones)

Glycoprotein-associated protein (113 clones)

Signal transduction-associated protein (148 clones)

Transcription-associated protein (233 clones)

Disease-associated protein (437 clones)

Enzyme or metabolism-associated protein (301 clones)

Cell division- or cell proliferation-associated protein (74 clones)

Cytoskeleton-associated protein (92 clones)

RNA synthesis-associated protein (280 clones)

Nuclear protein (352 clones)

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Protein synthesis- or transport-associated protein (112 clones)

Cellular defense-associated protein (23 clones)

Development- or growth-associated protein (23 clones)

[0060] It is also possible to predict the protein function by looking into the amino acid sequence for the motifs such

as the signal sequence, transmembrane region, nuclear translocation signal, glycosylation signal, phosphorylation site, Zinc finger motif, and SH3 domain. The programs, PSORT (Nakai K., and Kanehisa M. (1992) Genomics 14: 897-911), SOSUI (Hirokawa T. et al. (1998) Bioinformatics 14: 378-379) (Mitsui Information Developing Inc.), and MEMSAT (Jones D.T., Taylor W.R., and Thornton J.M. (1994) Biochemistry 33: 3038-3049) can be used to predict the existence of the signal sequence or transmembrane region. Alternatively, a partial amino acid sequence of the protein is fused with another protein such as GFP, the fusion protein is transfected into cultured cells, and the localization is analyzed to predict the function of the original protein.

[0061] Based on the determined nucleotide sequences of the full-length cDNAs obtained in the present invention, it is possible to predict more detailed functions of the proteins encoded by the cDNA clones, for example, by searching the databases such as GenBank, Swiss-Prot and UniGene for homologies of the cDNAs; or by searching the amino acid sequences deduced from the full-length cDNAs for signal sequences by using software programs such as PSORT, for transmembrane regions by using software programs such as SOSUI or for motifs by using software programs such as Pfam (http://www.expasy.ch/prosite/). As a matter of course, the functions are often predictable by using partial sequence information (preferably 300 nucleotides or more) instead of the full-length nucleotide sequences. However, the result of the prediction by using partial nucleotide sequence does not always agree with the result obtained by using full-length nucleotide sequence, and thus, it is needless to say that the prediction of function is preferably performed based on the full-length nucleotide sequences of the 4997 clones (see Example 18). The amino acid sequences deduced from the full-length nucleotide sequences were searched for functional domains by PSORT, SOSUI and Pfam. Prediction of functions of proteins encoded by the clones and the categorization thereof were performed based on these results obtained.

The following 798 clones were categorized into secretory and/or membrane proteins.

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                                              THYRO1000783,
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THYRO1001828,
               Y79AA1000258,
                               Y79AA1000420,
                               Y79AA1001177,
Y79AA1000976,
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Y79AA1001647.
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                               Y79AA1001874,
                                               Y79AA1002139,
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                                                                              Y79AA1002351,
Y79AA1002399, Y79AA1002416, MAMMA1002498, NT2RM4002287
[0062] The following 142 clones were categorized into glycoprotein-associated proteins.
HEMBA1000156, HEMBA1000518, HEMBA1000852, HEMBA1001071, HEMBA1001286,
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                              HEMBA1003071, HEMBA1003077, HEMBA1003281,
                                                                             HEMBA1003538,
HEMBA1003679,
              HEMBA1003866,
                              HEMBA1005576, HEMBA1005581, HEMBA1005699,
                                                                             HEMBA1006038,
HEMBA1006976,
              HEMBA1007301,
                              HEMBB1000317, HEMBB1000915, HEMBB1001871,
                                                                             HEMBB1001872,
HEMBB1002193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388, MAMMA1002329,
MAMMA1002428, MAMMA1002573, MAMMA1003150, NT2RM1000648, NT2RM1001115,
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NT2RM2000407,
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                              NT2RM2000490, NT2RM2001499,
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               NT2RM4002251,
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OVARC1000288,
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                              OVARC1001055,
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PLACE1000213, PLACE1000401, PLACE1002437, PLACE1002583,
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    THYRO1001287, Y79AA1001603, Y79AA1001874, MAMMA1002498
    [0063] The following 140 clones were categorized into signal transduction-associated proteins.
    HEMBA1000303, HEMBA1000369,
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                                                                                HEMBA1001019,
    HEMBA1001174, HEMBA1001822,
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                                                                                HEMBA1002341,
    HEMBA1002417, HEMBA1002768,
                                  HEMBA1003250, HEMBA1003291,
                                                                 HEMBA1003645,
                                                                                HEMBA1004286,
    HEMBA1005737, HEMBA1006130,
                                  HEMBA1006708, HEMBB1000083,
                                                                 HEMBB1000266,
                                                                                HEMBB1000632,
    HEMBB1000781, HEMBB1000831, HEMBB1002193, MAMMA1000173, MAMMA1001038,
    MAMMA1001198, MAMMA1002842, MAMMA1003057, NT2RM1000702, NT2RM1000772.
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                                   NT2RM4002013.
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                                                                 NT2RP5003477,
    OVARC1000437, OVARC1000556, OVARC1000649, OVARC1000945, OVARC1001200,
    OVARC1002182,
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    1000072, THYRO1000748, THYRO1001120, Y79AA1000328, Y79AA1002431, HEMBA1001247, NT2RM2001813,
    NT2RM4001454, NT2RP2005140, NT2RP2005293.
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    PLACE1003723, PLACE1005327, PLACE3000124.
    [0064] The following 321 clones were categorized into transcription -associated proteins.
    HEMBA1000158, HEMBA1000201, HEMBA1000216, HEMBA1000555, HEMBA1000561,
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    HEMBA1001077.
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    HEMBA1001809, HEMBA1001819.
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                                                                 HEMBA1002035,
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    HEMBA1002177.
                  HEMBA1002770.
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                                                 HEMBA1003408,
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    HEMBA1003662, HEMBA1003684,
                                  HEMBA1003760,
                                                 HEMBA1003953,
                                                                 HEMBA1004097,
                                                                                HEMBA1004321,
    HEMBA1004353, HEMBA1004389,
                                  HEMBA1004479,
                                                                 HEMBA1004973,
                                                                                HEMBA1005219,
                                                 HEMBA1004758,
    HEMBA1005359, HEMBA1005513,
                                  HEMBA1005528.
                                                 HEMBA1005548,
                                                                 HEMBA1005558.
                                                                                HEMBA1005931,
    HEMBA1006158, HEMBA1006248,
                                  HEMBA1006278.
                                                 HEMBA1006283,
                                                                 HEMBA1006347.
                                                                                HEMBA1006359.
    HEMBA1006559, HEMBA1006941,
                                  HEMBB1000789,
                                                 HEMBB1001011,
                                                                 HEMBB1001314,
                                                                                HEMBB1001482.
    HEMBB1001673, HEMBB1001749,
                                  HEMBB1001839.
                                                 HEMBB1001908.
                                                                 HEMBB1002134,
                                                                                HEMBB1002217,
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    HEMBB1002342, HEMBB1002607, MAMMA1000183, MAMMA1000388, MAMMA1001105, MAMMA1001222,
    MAMMA1001260, MAMMA1001627, MAMMA1001633, MAMMA1001743, MAMMA1001820, MAMMA1001837,
    MAMMA1002617, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000086, NT2RM1000746,
    NT2RM1000885,
                   NT2RM1000894,
                                  NT2RM1001092.
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    NT2RM2000740, NT2RM2001035,
                                  NT2RM2001105,
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    [0069] The following 296 clones presumably belong to DNA- and/or RNA-binding proteins. HEMBA1000158,
    HEMBA1000216,
                   HEMBA1000561, HEMBA1000591, HEMBA1000851, HEMBA1001088, HEMBA1001137,
    HEMBA1001405.
                   HEMBA1001510,
                                   HEMBA1001804, HEMBA1001809, HEMBA1001819, HEMBA1001847,
    HEMBA1001869.
                   HEMBA1002177.
                                   HEMBA1002935,
                                                  HEMBA1003408,
                                                                  HEMBA1003545, HEMBA1003568,
    HEMBA1003591,
                   HEMBA1003662,
                                   HEMBA1003684,
                                                                                  HEMBA1003805,
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    HEMBA1003953,
                                                                  HEMBA1004479,
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                                                  HEMBA1004389,
                                                                                  HEMBA1004669,
    HEMBA1004847,
                   HEMBA1004973,
                                   HEMBA1005202,
                                                                  HEMBA1005931,
                                                                                  HEMBA1006248,
                                                  HEMBA1005359,
    HEMBA1006278,
                   HEMBA1006283,
                                   HEMBA1006359,
                                                  HEMBA1006652,
                                                                  HEMBA1007087.
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40
    HEMBB1000264,
                   HEMBB1000789,
                                   HEMBB1001011,
                                                  HEMBB1001482,
                                                                  HEMBB1001736,
                                                                                  HEMBB1001749,
    HEMBB1001839, HEMBB1002217, MAMMA1000183, MAMMA1000284,
                                                                 MAMMA1000731, MAMMA1001105,
    MAMMA1001222, MAMMA1001260, MAMMA1001743, MAMMA1001837, MAMMA1002385, MAMMA1002617,
    MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000086, NT2RM1000539, NT2RM1000555,
    NT2RM1000666,
                   NT2RM1000691,
                                   NT2RM1000826.
                                                                                  NT2RM1001092,
                                                   NT2RM1000885.
                                                                  NT2RM1001059.
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    NT2RM4000996, NT2RM4001092, NT2RM4001140,
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                                   NT2RM4001483,
                                                   NT2RM4001592,
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                                                                                  NT2RM4001823,
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                                   NT2RM4001880,
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                                                                   NT2RP1000966,
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                                                   NT2RP2005763,
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                                    NT2RP3000603,
                                                    NT2RP3000624.
                                                                    NT2RP3000632.
                                                                                    NT2RP3000994.
     NT2RP3001057,
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                                    NT2RP3001120,
                                                    NT2RP3001150,
                                                                    NT2RP3001155,
                                                                                    NT2RP3001338,
     NT2RP3001398,
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                                    NT2RP3002399.
                                                    NT2RP3002876,
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                                                                                    NT2RP4000929,
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                                                    NT2RP4002078,
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                                    OVARC1000846,
                                                    OVARC1001232,
                                                                    OVARC1001271,
                                                                                   OVARC1001306,
     OVARC1001987,
                    OVARC1002112,
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                                    PLACE1003605,
                                                    PLACE1003704,
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    PLACE1007547,
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                                    PLACE1007688.
                                                                    PLACE1008132.
                                                    PLACE1007969.
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    PLACE1009246.
                    PLACE1009398,
                                    PLACE1009476.
                                                    PLACE1009622.
                                                                    PLACE1010053.
                                                                                    PLACE1010194,
    PLACE1010702.
                    PLACE1010870,
                                    PLACE1011056.
                                                    PLACE1011114,
                                                                    PLACE1011433.
                                                                                    PLACE2000427,
    PLACE3000009,
                    PLACE3000169.
                                    PLACE4000014.
                                                    PLACE4000156.
                                                                    PLACE4000192,
                                                                                    PLACE4000261,
    PLACE4000489,
                   SKNMC1000091,
                                   THYRO 1000085.
                                                    THYRO1000242.
                                                                    THYRO1000501,
                                                                                   THYRO1001100,
    THYRO1001189,
                    THYRO1001809.
                                     Y79AA1000037.
                                                    Y79AA1000349.
                                                                                    Y79AA1001211,
                                                                    Y79AA1000752,
    Y79AA1001312,
                    Y79AA1001391.
                                                    Y79AA1002103,
                                    Y79AA1001613,
                                                                    Y79AA1002472,
                                                                                    Y79AA1002482,
    HEMBA1004596, OVARC1000148, PLACE1003334, THYRO1001661,
    [0070] The following 66 clones presumably belong to the category of RNA synthesis-associated proteins.
    HEMBA1000591,
                   HEMBA1001579.
                                   HEMBA1003179, HEMBA1003591,
                                                                   HEMBA1006278,
                                                                                   HEMBB1000226,
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                    NT2RM1000852,
                                   NT2RM2000624,
                                                    NT2RM2001989,
                                                                   NT2RM2002100,
                                                                                   NT2RM4000191,
    NT2RM4001178.
                    NT2RM4002093,
                                    NT2RP1000035,
                                                    NT2RP1000272,
                                                                    NT2RP1000470,
                                                                                    NT2RP1001080,
    NT2RP2000153,
                    NT2RP2002928,
                                    NT2RP2003157,
                                                    NT2RP2004568,
                                                                    NT2RP2005126,
                                                                                    NT2RP2005436,
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                    NT2RP2005605,
                                    NT2RP2005776,
                                                    NT2RP2005942,
                                                                    NT2RP2006043,
                                                                                    NT2RP2006238,
    NT2RP3000361,
                    NT2RP3000397,
                                    NT2RP3001671,
                                                    NT2RP3004504,
                                                                    NT2RP4000078.
                                                                                    NT2RP4000111,
    NT2RP4000481,
                    NT2RP4000518.
                                    NT2RP4000614,
                                                    NT2RP4000929.
                                                                    NT2RP4001696.
                                                                                    NT2RP4002058.
    OVARC1001232,
                    OVARC1001577,
                                    PLACE1000406.
                                                    PLACE1000596.
                                                                    PLACE1000755,
                                                                                    PLACE1001739,
    PLACE1003704.
                    PLACE1003885,
                                    PLACE1004564.
                                                    PLACE1004814.
                                                                    PLACE1004902.
                                                                                    PLACE1005373.
    PLACE1005646,
                    PLACE1005876.
                                    PLACE1006196,
                                                    PLACE1006626.
                                                                    PLACE1006878,
                                                                                    PLACE1006917,
    PLACE1009476, PLACE1009925, PLACE1010194, PLACE1011114, THYRO1000121, Y79AA1001963,
    [0071] The following 184 clones presumably belong to protein synthesis- and/or protein transport-associated pro-
    HEMBA1000012.
                   HEMBA1000141, HEMBA1000592.
                                                   HEMBA1003617,
                                                                   HEMBA1003773,
                                                                                   HEMBA1004202,
                   HEMBA1004734, HEMBA1004847,
    HEMBA1004276,
                                                                   HEMBA1004930.
                                                                                   HEMBA1005047,
                                                   HEMBA1004929,
                   HEMBA1006031, HEMBA1006272,
    HEMBA1005202,
                                                   HEMBA1006474.
                                                                   HEMBA1006652.
                                                                                   HEMBA1006914.
                   HEMBA1007224, HEMBB1000915,
    HEMBA1006973,
                                                   HEMBB1001112.
                                                                   HEMBB1001137,
                                                                                   HEMBB1001736.
    HEMBB1001831, HEMBB1001915, MAMMA1000085, MAMMA1000734, MAMMA1001008,
                                                                                  MAMMA1002170,
    MAMMA1002219, MAMMA1002236, MAMMA1002619, NT2RM1000661, NT2RM1000833,
                                                                                   NT2RM2000092.
45
    NT2RM2000504,
                   NT2RM2000577,
                                   NT2RM2000821,
                                                   NT2RM2001201.
                                                                   NT2RM2001592.
                                                                                   NT2RM2001613.
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                                                   NT2RM2002055,
                                                                   NT2RM4000155,
                                                                                   NT2RM4000169,
    NT2RM4000344,
                   NT2RM4000356,
                                   NT2RM4000421,
                                                   NT2RM4000712.
                                                                   NT2RM4001054.
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                    NT2RP2001660.
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                                                    NT2RP2004791,
                                                                   NT2RP2005012,
                                                                                   NT2RP2005116,
    NT2RP2005360, NT2RP2005763, NT2RP2005784, NT2RP3000366,
    NT2RP3000759,
                    NT2RP3000968,
                                   NT2RP3001113,
                                                    NT2RP3001690,
                                                                   NT2RP3002045,
                                                                                   NT2RP3002151,
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                                                    NT2RP3003846,
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                                   OVARC1000564,
                                                   OVARC1000771,
                                                                  OVARC1000862,
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    OVARC1001180,
                    OVARC1001342,
                                    PLACE1000007,
                                                   PLACE1000061,
                                                                   PLACE1000081,
                                                                                  PLACE1000492,
    PLACE1000863,
                    PLACE1001092,
                                    PLACE1001748,
                                                   PLACE1002090,
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                                    PLACE1004743.
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    PLACE1006488,
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                                                                   PLACE1008273.
                                                                                  PLACE1008402,
                                   PLACE1009094,
    PLACE1008790.
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                                                                   PLACE1009477,
                                                                                  PLACE1009721,
    PLACE1009845,
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                                    PLACE1010547,
                                                   PLACE1011109,
                                                                   PLACE1011229,
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    PLACE1012031,
                    PLACE2000404,
                                    PLACE3000059,
                                                   PLACE3000121,
                                                                   PLACE4000269,
                                                                                  PLACE4000654,
    SKNMC1000011,
                    THYRO1000983,
                                    THYRO1001003,
                                                   THYRO1001313,
                                                                   Y79AA1000560,
                                                                                   Y79AA1000784,
    Y79AA1000968, Y79AA1001493, Y79AA1001875, Y79AA1002027, Y79AA1002209, HEMBA1006284,
    [0072] The following 130 clones presumably belong to cytoskeleton-associated proteins.
    HEMBA1000156, HEMBA1000168,
                                                                                  HEMBA1001651,
                                   HEMBA1000411, HEMBA1000588,
                                                                  HEMBA1001043,
    HEMBA1001661,
                    HEMBA1002102,
                                   HEMBA1002161.
                                                  HEMBA1002939.
                                                                  HEMBA1003235.
                                                                                  HEMBA1003581,
    HEMBA1004499,
                    HEMBA1004534.
                                   HEMBA1004697,
                                                  HEMBA1004929.
                                                                  HEMBA1004972.
                                                                                  HEMBA1005582,
    HEMBA1005595, HEMBA1006344.
                                   HEMBA1006737, HEMBB1001175,
                                                                  HEMBB1001282,
                                                                                  HEMBB1001562,
    HEMBB1001802, MAMMA1000824, MAMMA1001041, MAMMA1001576,
                                                                  MAMMA1001679, MAMMA1001735,
    MAMMA1002297, MAMMA1002351, MAMMA1002622, MAMMA1002637,
                                                                  MAMMA1003127, NT2RM1000850,
    NT2RM1000898, NT2RM2000030.
                                   NT2RM2000260.
                                                  NT2RM2000691.
                                                                  NT2RM2001324.
                                                                                  NT2RM4000169,
    NT2RM4000229, NT2RM4000515, NT2RM4001217, NT2RP1000202, NT2RP1000348, NT2RP1000460,
    NT2RP1000478,
                    NT2RP1001033,
                                   NT2RP1001294.
                                                   NT2RP1001302,
                                                                   NT2RP2000070.
                                                                                  NT2RP2000812,
    NT2RP2000814,
                    NT2RP2001168.
                                   NT2RP2001245.
                                                   NT2RP2001634.
                                                                   NT2RP2001900.
                                                                                  NT2RP2003307,
    NT2RP2003394,
                    NT2RP2004041.
                                   NT2RP2004242,
                                                                   NT2RP2004587,
                                                                                  NT2RP2004681.
                                                   NT2RP2004538,
    NT2RP2004732.
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                                                   NT2RP2005531,
                                                                   NT2RP2005712,
                                                                                  NT2RP2006275.
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                                   NT2RP3001216,
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                                                                   NT2RP3001272,
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                    NT2RP3001799.
                                                                                  NT2RP3003230.
                                   NT2RP3002688,
                                                   NT2RP3003061,
                                                                   NT2RP3003185,
    NT2RP3004569,
                    NT2RP3004578,
                                   NT2RP4001004,
                                                   NT2RP4001086,
                                                                   NT2RP4001256,
                                                                                  NT2RP4001567.
    NT2RP4001927,
                   OVARC1000001,
                                   OVARC1000106,
                                                   OVARC1000437,
                                                                  OVARC1000520,
                                                                                  OVARC1000679,
    OVARC1001731, OVARC1002050, PLACE1001104, PLACE1002571,
    PLACE1002591,
                    PLACE1002655,
                                   PLACE1002714,
                                                   PLACE1003625,
                                                                   PLACE1005287,
                                                                                   PLACE1006552,
    PLACE1007946,
                    PLACE1008426,
                                   PLACE1010148,
                                                   PLACE1010547,
                                                                   PLACE1010743,
                                                                                  PLACE1010896,
    PLACE1010960,
                    PLACE1011310.
                                   PLACE1011922,
                                                   PLACE2000216,
                                                                  PLACE2000274,
                                                                                  PLACE2000371,
    PLACE2000458.
                   PLACE3000145,
                                   PLACE3000416,
                                                   PLACE4000009.
                                                                  THYRO1000132,
                                                                                  THYRO1001405.
    THYRO1001458, Y79AA1000368, Y79AA1000794, Y79AA1000833, Y79AA1000962, Y79AA1002208,
35
    [0073]
          The following 54 clones presumably belong to cell division-associated and/or cell proliferation-associated
    proteins
    HEMBA1001019, HEMBA1001595, HEMBA1002363, HEMBA1002997, HEMBA1003136, HEMBA1003369,
    HEMBA1004131. HEMBA1004354.
                                   HEMBA1005621, HEMBB1000037, HEMBB1000264,
                                                                                 MAMMA1001768.
    MAMMA1002769. NT2RM1000354.
                                   NT2RM1000430, NT2RM1000874.
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    NT2RM2001896.
                   NT2RM2002145.
                                   NT2RM4000215,
                                                                   NT2RP1000163,
                                                                                  NT2RP1000333.
                                                   NT2RM4001714,
    NT2RP1000439,
                   NT2RP2000346,
                                   NT2RP2001397,
                                                   NT2RP2002595,
                                                                  NT2RP2003177,
                                                                                  NT2RP2003596,
    NT2RP2003912,
                                                   NT2RP2005520,
                   NT2RP2004396.
                                   NT2RP2005037.
                                                                  NT2RP2005669.
                                                                                  NT2RP2005835.
                                   NT2RP4000210,
    NT2RP3001730,
                   NT2RP3002081,
                                                                                  NT2RP4001634,
                                                   NT2RP4000415,
                                                                  NT2RP4001414,
    OVARC1000013, OVARC1000937, PLACE1001383,
                                                   PLACE1002433,
                                                                  PLACE1004316,
                                                                                  PLACE1005287,
    PLACE1008808, PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394
    [0074] The following 36 clones presumably belong to the category of embryogenesis- and/or development-associ-
    ated proteins:
    HEMBA1000518, HEMBA1001847,
                                   HEMBA1001869,
                                                  HEMBA1003545,
                                                                  HEMBA1004973,
                                                                                  HEMBB1002442,
    MAMMA1001837, NT2RM2001670,
                                   NT2RM4000046,
                                                   NT2RM4000531,
                                                                  NT2RM4001140,
                                                                                  NT2RM4001858.
    NT2RP2002078.
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                                   NT2RP2006436.
                                                   NT2RP3000603.
                                                                   NT2RP3000994,
                                                                                  NT2RP3001580,
    NT2RP3001708,
                   NT2RP3003071,
                                                                                  NT2RP4001567,
                                   NT2RP3004472.
                                                   NT2RP3004617.
                                                                   NT2RP4000246.
    OVARC1000304, OVARC1000746, PLACE1000793,
                                                   PLACE1002532,
                                                                   PLACE1003258,
                                                                                  PLACE1003625.
    PLACE1004460, PLACE1009622, PLACE4000558, THYRO1000085, Y79AA1001391, Y79AA1001692
    [0075] The following 30 clones presumably belong to cellular defense-associated proteins.
    HEMBA1000005, HEMBA1000531,
                                   HEMBA1003417,
                                                  HEMBA1006253,
                                                                  NT2RM4000354,
                                                                                  NT2RM4001880,
    NT2RP1000333.
                                   NT2RP2000006,
                   NT2RP1000493,
                                                   NT2RP2000045,
                                                                   NT2RP2000809,
                                                                                  NT2RP2001536,
    NT2RP2002464,
                                   NT2RP2005037,
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[0076] Although it is unclear whether or not 261 clones out of clones other than the above-mentioned clones belong to any of the above-described categories, these clones are predicted to have some functions, based on the homology search using their full-length sequences.

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[0077] In some cases, the predicted functions based on the partial sequences are different from those based on the full-length sequences. The reason is that a protein does not always belong solely to a single category of the above-described functional categories, and therefore, a protein may belong to two or more of the predicted functional categories. Besides, additional functions can be found for the clones classified into these functional categories by further analyses.

[0078] Since the protein encoded by clones of the invention contains full-length amino acid sequence, it is possible to analyze its biological activity, and its effect on cellular conditions such as cell proliferation and differentiation by expressing the protein as a recombinant protein using an appropriate expression system, injecting the recombinant into the cell, or raising a specific antibody against the protein.

[0079] If the protein is a secretory protein, membrane protein, or protein associated with glycoprotein, signal trans-

duction, or transcription, its biological activity can be analyzed by the methods in "Gene Transcription" (Hames B.D., and Higgins S.J. edit, (1993)), "Glycobiology" (Fukuda M., and Kobata A. edit, (1993)), "Growth Factors" (McKay I., and Leigh I. edit, (1993)), "Extracellular Matrix" (Haralson M.A., and Hassell J.R. edit, (1995), "Transcription Factors" (Latchman D.S. edit, (1993)), "Signal Transduction" (Milligans G. edit, (1992)), "Protein Phosphorylation" (Hardies G. D. edit, (1993), and "Ion Channels" (Ashley R.H. edit, (1995) featured in "The Practical Approach Series" (IRL PRESS), or "Signal Transduction Protocols" (Kendall D.A., and Hill S.J. edit, (1995), "Glycoprotein Analysis in Biomedicine" (Hounsell E.F. edit, (1993)), featured in "Method in Molecular Biology" (Humana Press).

[0080] As to a protein associated with a disease, it is possible to perform a functional analysis as described above, but also possible to analyze correlation between the expression or the activity of the protein and a certain disease by using a specific antibody that is obtained by using expressed protein. Alternatively, it is possible to utilize the database Online Mendelian Inheritance in Man (OMIM) (http://www.ncbi.nlm.nih.gov/Omim/), which is a database of human genes and diseases, to analyze the protein.

New information is constantly being deposited in the OMIM database. Therefore, it is possible for one skilled in the art to find a new relationship between a particular disease and a gene of the present invention in the most up-to-date database.

[0081] Also, as for a secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., search of the OMIM with the following keywords resulted in the finding that the proteins are associated with many diseases (the result of the OMIM search for secrete and membrane proteins is shown below). Also, association between proteins associated to signal transduction or transcription and diseases is reported in "Transcription Factor Research-1999" (Fujii, Tamura, Kageyama, and Satake edit, (1999) Jikken-Igaku Zoukan, Vol.17, No.3), and "Gene Medicine" ((1999) Vol.3, No.2). For example, in tumors, many proteins have been shown to play a role, including secretory proteins, membrane proteins, and proteins associated with signal transduction, glycoprotein, and transcription, and also proteins associated with metabolism, cytoskeleton, and cell cycle, as described in "Tumor Biology" (Matsubara S. (1992) Syoukabou Life Science series). Thus, besides the proteins associated with diseases, many proteins described above are also potentially associated with diseases, and thus useful as a target in the medicinal industry.

[0082] The result of the OMIM search for secretory and membrane proteins is shown below, in which the keywords,

- (1) secretion protein,
- (2) membrane protein,
- (3) channel, and

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(4) extracellular matrix were used.

[0083] Shown in the search result are only the accession numbers in the OMIM. Using the number, data showing the relationship between a disease and a gene or protein can be seen. The OMIM data has been renewed everyday.

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1) Secretion protein
         268 entries found, searching for "secretion protein"
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         2) Membrane protein
         1017 entries found, searching for "membrane protein"
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[0084] There are several methods for analyzing the expression levels of genes associated with diseases. Differences in gene expression levels between diseased and normal tissues are studied by the analytical methods, for example, Northern hybridization and differential display. Other examples include a method with high-density cDNA filter, a method with DNA microarray and methods with PCR amplification (Experimental Medicine, Vol.17, No. 8, 980-1056 (1999); Cell Engineering (additional volume) DNA Microarray and Advanced PCR Methods, Muramatsu & Naba (eds.), Shujunsya). The varying levels of gene expression between diseased tussues and normal tissues can be studied by any of these analytical methods. When explicit difference in the expression level is observed for a gene, it can be conclude that the gene is closely associated with a disease or disorder. Instead of diseased tissues, cultured cells can be used

for the assessment. Similarly, when gene expression is explicitly different between normal cells and cells reproducing disease-associated specific features, it can be concluded that the gene is closely associated with a disease or disorder. When the expression levels of genes are evidently varied during major cellular events (such as differentiation and apoptosis), the genes are involved in the cellular events and accordingly are candidates for disease- and/or disorder-associated genes. Further, genes exhibiting tissue-specific expression are genes playing important parts in the tissue functions and, therefore, can be candidates for genes associated with diseases and/or disorders affecting the tissues. [0085] For example, non-enzymic protein glycation reaction is believed to be a cause for a variety of chronic diabetic complications. Accordingly, genes of which expression levels are elevated or decreased in a glycated protein-dependent manner in the endothelial cells, are associated with diabetic complications caused by glycated proteins (Diabetes 1996, 45 (Suppl. 3), S67-S72; Diabetes, 1997, 46 (Suppl. 2), S19-S25).

The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)- $\alpha$  participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis. Many genes acting at the downstream of TNF- $\alpha$  and IL-1 $\beta$  among inflammation-associated cytokines have been previously identified. The respective stimulations are transduced through independent pathways of signaling cascade. There exists another signaling cascade for both stimulations, wherein NF- $\kappa$  B is a common transducing molecule shared by the two stimulations (J. Leukoc. Biol., 1994, 56(5): 542-547). It has also been revealed that many inflammation-associated genes, including IL-2, IL-6 and G-CSF, are varied in the expression levels thereof in response to the signal through the common pathway (Trend Genet. 1999, 15(6): 229-235). It is assumed that genes of which expression levels are varied in response to the stimulation of TNF- $\alpha$  or IL-1 $\beta$  also participate in inflammation.

[0086] Ultraviolet radiation damage has been recognized as a risk factor for skin cancers, etc. (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with ultraviolet radiation damage of skin. In addition, genes associated with neural differentiation can be candidates for genes responsible for neurological diseases as well as candidates for genes usable for treating the diseases.

[0087] Clones exhibiting differences in the expression levels thereof can be selected by using gene expression analysis. The selection comprises, for example; analyzing cDNA clones by using high-density cDNA filter; and statistically treating the multiple signal values (signal values of radioisotope in the labeled probes or values obtained by measuring fluorescence intensities emitted from the fluorescent labels) for the respective clones by two-sample t-test, where the signal values are determined by multiple experiments of hybridization. The clones of interest are selectable based on the statistically significant differences in the signal distribution at p<0.05. However, selectable clones with significant difference in the expression levels thereof may be changed depending on the partial modification of statistical treatment. For example, the clones may be selected by conducting statistical treatment with two-sample t-test at p<0.01; or genes exhibiting more explicit differences in the expression levels thereof can be selected by performing statistical treatment with a pre-determined cut-off value for the significant signal difference. An alternative method is that the expression levels are simply compared with each other, and then, the clones of interest are selected based on the ratio of the expression levels thereof.

[0088] Clones that vary in their expression levels can also be selected by comparing the expression levels by PCR analysis, for example, by using the method of determining the band intensities representing the amounts of PCR products with ethidium bromide staining; the method of determining the values of radioisotope signals or fluorescence intensities of the PCR products when radiolabeled or fluorescent dye-labeled primers, respectively, are used in PCR amplification; or the method of determining the values of radioisotope signals or fluorescence intensities of the probes hybridized to the PCR products when radiolabeled or fluorescent dye-labeled probes, respectively, are used in the hybridization. If the expression level ratios obtained in multiple PCR experiments are constantly at least 2-fold, such a clone can be judged to vary in its expression level. When the ratios are several-fold or not less than 10-fold, the clone can be selected as a gene exhibiting the explicit difference in its expression level.

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[0089] A survey of genes of which expression levels are varied specifically to the glycated protein in the endothelial cells has revealed genes with elevated expression levels, HEMBA1003958, HEMBA1004850, MAMMA1001256, MAMMA1002132, PLACE2000411 and PLACE3000119. On the other hand, a gene of which expression level is decreased specifically to the glycated protein is MAMMA1001783. These clones are genes associated with diabetes.

[0090] A survey of genes of which expression levels are varied in response to TNF- $\alpha$  (Tumor Necrosis Factor-alpha) in the primary cell culture of synovial tissue has revealed the following clones with elevated expression levels in the presence of TNF- $\alpha$ :

HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000046, HEMBA1000076, HEMBA1000111, HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231, HEMBA1000243, HEMBA1000280,

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HEMBA1000282, HEMBA1000304, HEMBA1000307,
                                           HEMBA1000327, HEMBA1000356, HEMBA1000376,
HEMBA1000387, HEMBA1000390, HEMBA1000418,
                                           HEMBA1000460.
                                                          HEMBA1000491.
                                                                        HEMBA1000501.
HEMBA1000518,
              HEMBA1000519,
                            HEMBA1000520,
                                           HEMBA1000531,
                                                          HEMBA1000534,
                                                                         HEMBA1000542,
HEMBA1000545,
              HEMBA1000591,
                            HEMBA1000592.
                                           HEMBA1000594,
                                                          HEMBA1000636,
                                                                        HEMBA1000655,
HEMBA1000657, HEMBA1000673,
                             HEMBA1000682,
                                           HEMBA1000686,
                                                          HEMBA1000722,
                                                                        HEMBA1000726.
HEMBA1000827, HEMBA1000870,
                             HEMBA1000918,
                                           HEMBA1000971,
                                                          HEMBA1000974, HEMBA1000986,
HEMBA1001019, HEMBA1001043,
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HEMBA1001080, HEMBA1001109,
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                                           HEMBA1002241,
                                                          HEMBA1002337,
                                                                         HEMBA1002363,
HEMBA1002389, HEMBA1002458,
                             HEMBA1002460,
                                           HEMBA1002469,
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HEMBA1002659, HEMBA1002750,
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                                                                        HEMBA1002816,
HEMBA1002818, HEMBA1002850,
                             HEMBA1002863.
                                           HEMBA1003021.
                                                          HEMBA1003033.
                                                                         HEMBA1003078,
HEMBA1003166, HEMBA1003202,
                             HEMBA1003204.
                                           HEMBA1003229.
                                                          HEMBA1003235.
                                                                         HEMBA1003276.
HEMBA1003286, HEMBA1003296.
                             HEMBA1003370.
                                           HEMBA1003376.
                                                          HEMBA1003403.
                                                                         HEMBA1003418.
HEMBA1003433. HEMBA1003447.
                             HEMBA1003560.
                                           HEMBA1003569.
                                                          HEMBA1003571.
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[0092] These are rheumatoid arthritis-associated clones. [0093] A survey of genes of which expression levels are varied in primary cultured skin fibroblast cells exposed to ultraviolet light has revealed the following clones with elevated expression levels by ultraviolet radiation: HEMBA1000542, HEMBA1001808, HEMBA1002177, HEMBA1003314, MAMMA1001874, NT2RM2001100, NT2RP2005732, NT2RP3000592, NT2RP4000657, OVARC 1000004, OVARC1001092, OVARC1001342, PLACE1002816, NT2RM4001002, NT2RM4001813, NT2RM4002266, NT2RP2001174, NT2RP2001196, NT2RP3001216, NT2RP2005358. NT2RP3000690, NT2RP3003464, PLACE1006382, THYRO1000070, THYRO1001100, Y79AA1000342 [0094] On the other hand, the expression levels of the following clones were decreased by ultraviolet radiation: HEMBA1000005, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000168, HEMBA1000231, HEMBA1000304, HEMBA1000307, HEMBA1000333, HEMBA1000366, HEMBA1000369, HEMBA1000390, HEMBA1000396, HEMBA1000418, HEMBA1000434, HEMBA1000464, HEMBA1000469, HEMBA1000490, HEMBA1000504, HEMBA1000505, HEMBA1000557, HEMBA1000657, HEMBA1000673, HEMBA1000682, HEMBA1000686, HEMBA1000727, HEMBA1000752, HEMBA1000851, HEMBA1000852, HEMBA1000870, HEMBA1000872, HEMBA1001085, HEMBA1001121, HEMBA1001133, HEMBA1001235. HEMBA1001265, HEMBA1001281, HEMBA1001289, HEMBA1001299, HEMBA1001303. HEMBA1001310. HEMBA1001323, HEMBA1001595, HEMBA1001620, HEMBA1001678. HEMBA1001640. HEMBA1001712, HEMBA1001835. HEMBA1001950, HEMBA1001987. HEMBA1002253. HEMBA1002321, HEMBA1002341, HEMBA1002419. HEMBA1002679, HEMBA1002728. HEMBA1002818, HEMBA1002935, HEMBA1002999. HEMBA1003034, HEMBA1003071, HEMBA1003175, HEMBA1003098. HEMBA1003142, HEMBA1003202, HEMBA1003212, HEMBA1003276, HEMBA1003220, HEMBA1003373. HEMBA1003417, HEMBA1003447, HEMBA1003528. HEMBA1003885, HEMBA1003684, HEMBA1003799, HEMBA1003989. HEMBA1004011, HEMBA1004055, HEMBA1004133, HEMBA1004225, HEMBA1004272, HEMBA1004353, HEMBA1004631, HEMBA1004669, HEMBA1004705. HEMBA1004753, HEMBA1004776, HEMBA1004900, HEMBA1004803, HEMBA1004816, HEMBA1005047, HEMBA1005079, HEMBA1005101, HEMBA1005149, HEMBA1005152, HEMBA1005202, HEMBA1005314, HEMBA1005372, HEMBA1005511, HEMBA1005513, HEMBA1005518, HEMBA1005570, HEMBA1005577, HEMBA1005581, HEMBA1005588, HEMBA1005609, HEMBA1005632, HEMBA1005853, HEMBA1006031, HEMBA1006035, HEMBA1006485, HEMBA1006486, HEMBA1006502, HEMBA1006696, HEMBA1006789, HEMBA1006796, HEMBA1007085, HEMBA1007224, HEMBA1007301, HEMBA1007319, HEMBA1007341, HEMBA1007342, HEMBB1000036, HEMBB1000037, HEMBB1000217, HEMBB1000266, HEMBB1000317, HEMBB1000336, HEMBB1000354, HEMBB1000369, HEMBB1000399, HEMBB1000434, HEMBB1000438, HEMBB1000592, HEMBB1000673, HEMBB1000789, HEMBB1000810, HEMBB1000883, HEMBB1000887, HEMBB1001105, HEMBB1001182, HEMBB1001242. HEMBB1001267, HEMBB1001424, HEMBB1001464, HEMBB1001531, HEMBB1001618. HEMBB1001996, HEMBB1002092, HEMBB1002139. 35 HEMBB1002142, HEMBB1002190. HEMBB1002453. HEMBB1002520, HEMBB1002550, HEMBB1002556. HEMBB1002600. HEMBB1002664. MAMMA1000009. MAMMA1000055. MAMMA1000069, MAMMA1000133, MAMMA1000287, MAMMA1000171, MAMMA1000173, MAMMA1000416, MAMMA1000585, MAMMA1000713. MAMMA1000760, MAMMA1000798, MAMMA1000831. MAMMA1000875, MAMMA1000876, MAMMA1000877, MAMMA1000906, MAMMA1000931, MAMMA1000962. MAMMA1001243. MAMMA1001133, MAMMA1001139, MAMMA1001271, MAMMA1001274, MAMMA1001298, MAMMA1001606, MAMMA1001630, MAMMA1001670, MAMMA1001743, MAMMA1001751, MAMMA1002140, MAMMA1002145, MAMMA1002158, MAMMA1002170, MAMMA1002236, MAMMA1002311, MAMMA1002498, MAMMA1002754. MAMMA1002780. MAMMA1002820, MAMMA1002843, MAMMA1002844, MAMMA1002871, MAMMA1003047, NT2RM1000037, NT2RM1000039, NT2RM1000080, NT2RM1000086, NT2RM1000341, NT2RM1000499, NT2RM1000669, NT2RM1000746, 45 NT2RM1000781, NT2RM1000885, NT2RM1000905, NT2RM1000962, NT2RM2000239, NT2RM2000260, NT2RM2000371, NT2RM2000639, NT2RM2000649, NT2RM2000735, NT2RM2000821, NT2RM2000984, NT2RM2001035, NT2RM2001065, NT2RM2001105, NT2RM2001177, NT2RM2001194, NT2RM2001196, NT2RM2001243, NT2RM2001256, NT2RM2001424, NT2RM2001588, NT2RM2001635, NT2RM2001648, NT2RM2001652, NT2RM2001668, NT2RM2001706, NT2RM2001727, NT2RM2001730, NT2RM2001743, NT2RM2001753, NT2RM2001760, NT2RM2001771, NT2RM2001785, NT2RM2001855. NT2RM2001800, NT2RM2001896, NT2RM2001997, NT2RM2002030, NT2RM2002049, NT2RM2002091, NT2RM2002142, NT2RM2002145, NT2RM2002178. NT2RM2002580, NT2RM4000215, NT2RM4000368, NT2RM4000344, NT2RM4000421. NT2RM4000425, NT2RM4000457, NT2RM4000496, NT2RM4000515, NT2RM4000712, NT2RM4000787. NT2RM4000813, NT2RM4000820, NT2RM4000852, NT2RM4000996, NT2RM4000950, NT2RM4001016, NT2RM4001047, NT2RM4001054, NT2RM4001140, NT2RM4001151, NT2RM4001187, NT2RM4001204, NT2RM4001258, NT2RM4001437, NT2RM4001454, NT2RM4001489, NT2RM4001605, NT2RM4001611, NT2RM4001666, NT2RM4001710, NT2RM4001714, NT2RM4001715, NT2RM4001731, NT2RM4001741, NT2RM4001746, NT2RM4001856, NT2RM4001938, NT2RM4001940, NT2RM4001984,

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     [0095] These clones are associated with ultraviolet radiation damage.
     [0096] A survey of genes of which expression levels are varied in response to the stimulation for inducing cell dif-
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    [0097] On the other hand, clones of which expression levels decreased by RA are as follows:
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    OVARC1000431, OVARC1001942,
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    PLACE1007257, PLACE1010624, Y79AA1000346
    [0098] Clones of which expression levels increase by RA/inhibitor are as follows:
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    Y79AA1001827, Y79AA1001875, Y79AA1002027, Y79AA1002211, Y79AA1002234, Y79AA1002258
    [0099] On the other hand, clones of which expression levels decrease by RA/inhibitor are as follows:
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                                                                    PLACE1011219,
                                                                                    THYRO1000270,
    Y79AA1000346, Y79AA1001541
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[0100] These clones are also associated with neural differentiation and, therefore, are candidates for genes associated with neurological diseases.

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[0101] For example, if the protein encoded by the cDNA of the present invention is a regulatory factor of cellular conditions such as growth and differentiation, it can be used for developing medicines as follows. The protein or antibody provided by the invention is injected into a certain kind of cells by microinjection. Then, using the cells, it is possible to screen low molecular weight compounds by measuring the change in the cellular conditions, or the activation or inhibition of a particular gene. The screening can be performed as follows. First, the protein is expressed and purified as recombinant. The purified protein is microinjected into cells such as various cell lines, or primary culture cells, and the cellular change such as growth and differentiation can be examined. Alternatively, the induction of genes whose expression is known to be associated with a particular change of cellular conditions may be detected by the amount of mRNA or protein. Or, the amount of intracellular molecules (low molecular weight compounds, etc.) that is changed by the function of the gene product (protein) which is known to be associated with a particular change of cellular conditions may be detected. The compounds to be screened (both low and high molecular compounds are acceptable) can be added to the culture media and assessed for their activity by measuring the change of the cellular conditions. Instead of microinjection, cell lines introduced with the gene obtained in the invention can be used for the screening. If the gene product is turn out to be associated with a particular change in the cellular conditions, the change of the product can be used as a measurement for screening. Once a compound is screened out which can activate or inhibit

the function of the protein of the invention, it can be applied for developing medicines.

**[0102]** If the protein encoded by the cDNA of the present invention is a secretory protein, membrane protein, or protein associated with signal transduction, glycoprotein, transcription, or diseases, it can be used in functional assays for developing medicines.

[0103] In case of a membrane protein, it is most likely to be a protein that functions as a receptor or ligand on the cell surface. Therefore, it is possible to reveal a new relationship between a ligand and receptor by screening the membrane protein of the invention based on the binding activity with the known ligand or receptor. Screening can be performed according to the known methods.

[0104] For example, a ligand against the protein of the invention can be screened in the following manner. Namely, a ligand that binds to a specific protein can be screened by a method comprising the steps of: (a) contacting a test sample with the protein of the invention or a partial peptide thereof, or cells expressing these, and (b) selecting a test sample that binds to said protein, said partial peptide, or said cells.

[0105] On the other hand, for example, screening using cells expressing the protein of the present invention that is a receptor protein can also be performed as follows. It is possible to screen receptors that is capable of binding to a specific protein by using procedures (a) attaching the sample cells to the protein of the invention or its partial peptide, and (b) selecting cells that can bind to the said protein or its partial peptide.

[0106] In a following screening as an example, first the protein of the invention is expressed, and the recombinant protein is purified. Next, the purified protein is labeled, binding assay is performed using a various cell lines or primary cultured cells, and cells that are expressing a receptor are selected (Growth and differentiation factors and their receptors, Shin-Seikagaku Jikken Kouza Vol.7 (1991) Honjyo, Arai, Taniguchi, and Muramatsu edit, p203-236, Tokyo-Kagaku-Doujin). A protein of the invention can be labeled with RI such as <sup>125</sup>I, and enzyme (alkaline phosphatase etc.). Alternatively, a protein of the invention may be used without labeling and then detected by using a labeled antibody against the protein. The cells that are selected by the above screening methods, which express a receptor of the protein of the invention, can be used for the further screening of an agonists or antagonists of the said receptor.

[0107] Once the ligand binding to the protein of the invention, the receptor of the protein of the invention or the cells expressing the receptor are obtained by screening, it is possible to screen a compound that binds to the ligand and receptor. Also it is possible to screen a compound that can inhibit both bindings (agonists or antagonists of the receptor, for example) by utilizing the binding activities.

[0108] When the protein of the invention is a receptor, the screening method comprises the steps of (a) contacting the protein of the invention or cells expressing the protein of the invention with the ligand, in the presence of a test sample, (b) detecting the binding activity between said protein or cells expressing said protein and the ligand, and (c) selecting a compound that reduces said binding activity when compared to the activity in the absence of the test sample. Furthermore, when the protein of the invention is a ligand, the screening method comprises the steps of (a) contacting the protein of the invention with its receptor or cells expressing the receptor in the presence of samples, (b) detecting the binding activity between the protein and its receptor or the cells expressing the receptor, and (c) selecting a compound that can potentially reduce the binding activity compared to the activity in the absence of the sample.

[0109] Samples to screen include cell extracts, expressed products from a gene library, synthesized low molecular compound, synthesized peptide, and natural compounds, for example, but are not construed to be listed here. A compound that is isolated by the above screening using a binding activity of the protein of the invention can also be used as a sample.

[0110] A compound isolated by the screening may be a candidate to be an agonist or an antagonist of the receptor of the protein. By utilizing an assay that monitors a change in the intracellular signaling such as phosphorylation which results from reduction of the binding between the protein and its receptor, it is possible to identify whether the obtained compound is an agonist or antagonist of the receptor. Also, the compound may be a candidate of a molecule that can inhibit the interaction between the protein and its associated proteins (including a receptor) in vivo. Such compounds can be used for developing drugs for precaution or cures of a disease with which the protein is associated.

[0111] Secretory proteins may regulate cellular conditions such as growth and differentiation. It is possible to find out a novel factor that regulates cellular conditions by adding the secretory protein of the invention to a certain kind of cell, and performing a screening by utilizing the cellular changes in growth or differentiation, or activation of a particular gene.

[0112] The screening can be performed, for example, as follows. First, the protein of the invention is expressed and purified in a recombinant form. Then, the purified protein is added to a various kind of cell lines or primary cultured cells, and the change in the cell growth and differentiation is monitored. The induction of a particular gene that is known to be involved in a certain cellular change is detected by the amounts of mRNA and protein. Alternatively, the amount of an intracellular molecule (low-molecular-weight compounds, etc.) that is changed by the function of a gene product (protein) that is known to function in a certain cellular change is used for the detection.

[0113] Once the screening reveals that the protein of the invention can regulate cellular conditions or the functions, it is possible to apply the protein as a pharmaceutical and diagnostic medicine for associated diseases by itself or by

altering a part of it into an appropriate composition.

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[0114] As is above described for membrane proteins, the secretory protein provided by the invention may be used to explore a novel ligand-receptor interaction using a screening based on the binding activity to a known ligand or receptor. A similar method can be used to identify an agonist or antagonist. The resulting compounds obtained by the methods can be a candidate of a compound that can inhibit the interaction between the protein of the invention and an interacting molecule (including a receptor). The compounds may be able to use as a preventive, therapeutic, and diagnostic medicine for the diseases, in which the protein may play a certain role.

[0115] Proteins associated with signal transduction or transcription may be a factor that affects a certain protein or gene in response to intracellular/extracellular stimuli. It is possible to find out a novel factor that can affect a protein or gene by expressing the protein provided by the invention in a certain types of cells, and performing a screening utilizing the activation of a certain intracellular protein or gene.

[0116] The screening may be performed as follows. First, a transformed cell line expressing the protein is obtained. Then, the transformed cell line and the untransformed original cell line are compared for the changes in the expression of a certain gene by detecting the amount of its mRNA or protein. Alternatively, the amount of an intracellular molecule (low molecular weight compounds) that is changed by the function of a certain gene product (protein) may be used for the detection. Furthermore, the change of the expression of a certain gene can be detected by introducing a fusion gene that comprises a regulatory region of the gene and a marker gene (luciferase, beta-galactosidase, etc.) into a cell, expressing the protein provided by the invention into the cell, and estimating the activity of a marker gene product (protein).

[0117] If the protein or gene of the invention is associated with diseases, it is possible to screen a gene or compound that can regulate its expression and/or activity either directly or indirectly by utilizing the protein of the present invention.

[0118] For example, the protein of the invention is expressed and purified as a recombinant protein. Then, the protein or gene that interacts with the protein of the invention is purified, and screened based on the binding. Alternatively, the screening can be performed by adding with a compound of a candidate of the inhibitor added in advance and monitoring the change of binding activity. In another method, a transcription regulatory region locating in the 5'-upstream of the gene encoding the protein of the invention that is capable of regulating the expression of other genes is obtained, and fused with a marker gene. The fusion is introduced into a cell, and the cell is added with compounds to explore a regulatory factor of the expression of the said gene.

[0119] The compound obtained by the screening can be used for developing pharmaceutical and diagnostic medicines for the diseases with which the protein of the present invention is associated. Similarly, if the regulatory factor obtained in the screening is turn out to be a protein, compounds that can newly affect the expression or activity of the protein may be used as a medicine for the diseases with which the protein of the invention is associated.

**[0120]** If the protein of the invention has an enzymatic activity, regardless as to whether it is a secretory protein, membrane protein, or proteins associated with signal transduction, glycoprotein, transcription, or diseases, a screening may be performed by adding a compound to the protein of the invention and monitoring the change of the compound. The enzymatic activity may also be utilized to screen a compound that can inhibit the activity of the protein.

[0121] In a screening given as an example, the protein of the invention is expressed and the recombinant protein is purified. Then, compounds are contacted with the purified protein, and the amount of the compound and the reaction products is examined. Alternatively, compounds that are candidates of an inhibitor are pretreated, then a compound (substrate) that can react with the purified protein is added, and the amount of the substrate and the reaction products is examined.

[0122] The compounds obtained in the screening may be used as a medicine for diseases with which the protein of the invention is associated. Also they can be applied for tests that examine whether the protein of the invention functions normally *in vivo*.

[0123] Whether the secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein of the present invention is a novel protein associated with diseases or not is determined in another method than described above, by obtaining a specific antibody against the protein of the invention, and examining the relationship between the expression or activity of the protein and a certain disease. In an alternative way, it may be analyzed referred to the methods in "Molecular Diagnosis of Genetic Diseases" (Elles R. edit, (1996) in the series of "Method in Molecular Biology" (Humana Press).

**[0124]** Proteins associated with diseases are targets of screening as mentioned, and thus are very useful in developing drugs which regulate their expression and activity. Also, the proteins are useful in the medicinal industry as a diagnostic marker of the associated disease or a target of gene therapy.

[0125] Compounds isolated as mentioned above can be administered patients as it is, or after formulated into a pharmaceutical composition according to the known methods. For example, a pharmaceutically acceptable carrier or vehicle, specifically sterilized water, saline, plant oil, emulsifier, or suspending agent can be mixed with the compounds appropriately. The pharmaceutical compositions can be administered to patients by a method known to those skilled in the art, such as intraarterial, intravenous, or subcutaneous injections. The dosage may vary depending on the weight

or age of a patient, or the method of administration, but those skilled in the art can choose an appropriate dosage properly. If the compound is encoded by DNA, the DNA can be cloned into a vector for gene therapy, and used for gene therapy. The dosage of the DNA and the method of its administration may vary depending on the weight or age of a patient, or the symptoms, but those skilled in the art can choose properly.

[0126] The present invention further relates to databases comprising at least a sequence of polynucleotide and/or protein, or a medium recorded in such databases, selected from the sequence data of the nucleotide and/or the amino acids indicated in Table 350 and Table 351.

The term "database" means a set of accumulated information as machine-searchable and readable information of nucleotide sequence. The databases of the present invention comprise at least one of the novel nucleotide sequences of polynucleotides provided by the present invention. The databases of the present invention can consist of only the sequence data of the novel polynucleotides provided by the present invention or can comprise other information on nucleotide sequences of known full-length cDNAs or ESTs. The databases of the present invention can be comprised of not only the information on the nucleotide sequences but also the information on the gene functions revealed by the present invention. Additional information such as names of DNA clones carrying the full-length cDNAs can be recorded or linked together with the sequence data in the databases.

[0127] The database of the present invention is useful for gaining complete gene sequence information from partial sequence information of a gene of interest. The database of the present invention comprises nucleotide sequence information of full-length cDNAs. Consequently, by comparing the information in this database with the nucleotide sequence of a partial gene fragment yielded by differential display method or subtraction method, the information on the full-length nucleotide sequence of interest can be gained from the sequence of the partial fragment as a starting clue.

[0128] The sequence information of the full-length cDNAs constituting the database of the present invention contains not only the information on the complete sequences but also extra information on expression frequency of the genes as well as homology of the genes to known genes and known proteins. Thus the extra information facilitates rapid functional analyses of partial gene fragments. Further, the information on human genes is accumulated in the database of the present invention, and therefore, the database is useful for isolating a human homologue of a gene originating from other species. The human homologue can be isolated based on the nucleotide sequence of the gene from the original species.

[0129] At present, information on a wide variety of gene fragments can be obtained by differential display method and subtraction method. In general, these gene fragments are utilized as tools for isolating the full-length sequences thereof. When the gene fragment corresponds to an already-known gene, the full-length sequence is easily obtained by comparing the partial sequence with the information in known databases. However, when there exists no information corresponding to the partial sequence of interest in the known databases, cDNA cloning should be carried out for the full-length cDNA. It is often difficult to obtain the full-length nucleotide sequence using the partial sequence information as an initial clue. If the full-length of the gene is not available, the amino acid sequence of the protein encoded by the gene remains unidentified. Thus the database of the present invention can contribute to the identification of full-length cDNAs corresponding to gene fragments, which cannot be revealed by using databases of known genes.

[0130] The present invention has provided 5602 novel full-length cDNA clones, and primers for synthesizing the cDNA. As has not yet proceeded the isolation of full-length cDNA within the human, the invention has great significance. The full-length cDNA clones contain the translation initiation site, and thus provide a useful information for analysis of protein functions.

**[0131]** The cDNA clones are assumed to encode proteins such as secretory proteins, membrane proteins, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., which have important functions in vivo, and also predicted to be associated with many diseases. The genes and proteins associated with diseases are useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as a target of gene therapy.

[0132] The invention is illustrated more specifically with reference to the following examples, but is not to be construed as being limited thereto.

# **EXAMPLE 1**

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Construction of a cDNA library by the oligo-capping method.

[0133] The NT-2 neuron progenitor cells (Stratagene), a teratocarcinoma cell line from human embryo testis, which can differentiate into neurons by the treatment with retinoic acid were used.

The NT-2 cells were cultured according to the manufacturer's instructions as follows.

- (1) NT-2 cells were cultured without induction by retinoic acid treatment (NT2RM1, NT2RM2, NT2RM4).
- (2) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 48 hours (NT2RP1).

(3) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 2 weeks (NT2RP2, NT2RP3, NT2RP4, NT2RP5).

[0134] Also, the human neuroblastoma cell line SK-N-MC (ATCC HTB-10) (SKNMC1), and human retinoblastoma cell line Y79 (ATCC HTB-18) (Y79AA1) were cultured according to the culture conditions described in the ATCC catalogue (http://www.atcc.org/). The cells were harvested separately, and mRNA was extracted from each cell by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose. [0135] Similarly, human placenta (PLACE1, PLACE2, PLACE3, PLACE4), human ovary cancer tissue (OVARC1), tissues from human embryo at 10 weeks, which is enriched with head (HEMBA1), or body (HEMBB1), human mammary gland (MAMMA1), human thyroid gland (THYRO1), and primary cultured cells of human blood vessel endothelium (VESEN1) were used to extract mRNA by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose.

[0136] Each poly(A)\*RNA was used to construct a cDNA library by the oligo-capping method (Maruyama M. and Sugano S. (1994) Gene, 138: 171-174). Using the Oligo-cap linker (SEQ ID NO: 10464) and the Oligo-dT primer (SEQ ID NO: 10465), bacterial alkaline phosphatase (BAP) treatment, tobacco acid phosphatase (TAP) treatment, RNA ligation, the first strand cDNA synthesis, and removal of RNA were performed as described in the reference (Suzuki and Kanno (1996) Protein Nucleic acid and Enzyme, 41: 197-201; Suzuki Y. et al. (1997) Gene, 200: 149-156), Next, 5'- and 3'-PCR primers (SEQ ID NO: 10466, and 10467, respectively) were used for performing PCR to convert the cDNA into double stranded cDNA, which was then digested with Sfil. Then, the DrallI-cleaved pUC19FL3 vector (Figure 1; for NT2RM1, and NT2RP1), or the DrallI-cleaved pME18SFL3 (Figure 1) (GenBank AB009864, expression vector; for NT2RM2, NT2RM4, NT2RP2, NT2RP3, NT2RP4, NT2RP5, SKNMC1, Y79AA1, PLACE1, PLACE2, PLACE3, PLACE4, OVARC1, HEMBA1, HEMBB1, MAMMA1, THYRO1, and VESEN1) was used for cloning the cDNA in a unidirectional manner, and cDNA libraries were obtained. The nucleotide sequence of the 5'- and 3'- ends of the cDNA clones was analyzed with a DNA sequencer (ABI PRISM 377, PE Biosystems) after sequencing reactions were performed with the DNA sequencing reagents (Dye Terminator Cycle Sequencing FS Ready Reaction Kit, dRhodamine Terminator Cycle Sequencing FS Ready Reaction Kit, or BigDye Terminator Cycle Sequencing FS Ready Reaction Kit, PE Biosystems), according to the instructions. The data were compiled into a database.

[0137] The full-length-enriched cDNA libraries except those for NT2RM1 and NT2RP1 were constructed using eukaryotic expression vector pME18SFL3. The vector contains SRα promoter and SV40 small t intron in the upstream of the cloning site, and SV40 polyA added signal sequence site in the downstream. As the cloning site of pME18SFL3 has asymmetrical DrallI sites, and the ends of cDNA fragments contain Sfil sites complementary to the DrallI sites, the cloned cDNA fragments can be inserted into the downstream of the SRa promoter unidirectionally. Therefore, clones containing full-length cDNA can be expressed transiently by introducing the obtained plasmid directly into COS cells. Thus, the clones can be analyzed very easily in terms of the proteins that are the gene products of the clones, or in terms of the biological activities of the proteins.

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[0138] Herein, the cDNA libraries and the name of each clone are related as shown in Table 3. Therein, "xxxxxx" represents the clone number of six digits. Thus, the sequences are named by the library name, the clone number plus F- for the 5'-end, or R- for the 3'-end.

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Table 3

5	library: clone	5'-end sequence	3'-end sequence
10	NT2RM1: NT2RM1xxxxxx NT2RP1:	F-NT2RM1xxxxxx	
	NT2RP1xxxxxx NT2RM2:	F-NT2RP1xxxxxx	
15	NT2RM2xxxxxx NT2RM4:	F-NT2RM2xxxxxx	R-NT2RM2xxxxxx
	NT2RM4xxxxxx	F-NT2RM4xxxxxx	R-NT2RM4xxxxxx
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	NT2RP2:		
	NT2RP2xxxxxx	F-NT2RP2xxxxxx	R-NT2RP2xxxxxx
5	NT2RP3:		
	NT2RP3xxxxxx	F-NT2RP3xxxxxx	R-NT2RP3xxxxxx
	NT2RP4:		
	NT2RP4xxxxxx	F-NT2RP4xxxxxx	R-NT2RP4xxxxxx
10	NT2RP5:		
	NT2RP5xxxxxx	F-NT2RP5xxxxxx	R-NT2RP5xxxxxx
	SKNMC1:		
15	SKNMC1xxxxxx	F-SKNMC1xxxxxx	R-SKNMC1xxxxxx
15	Y79AA1:		
	Y79AA1xxxxxx	F-Y79AA1xxxxxx	R-Y79AA1xxxxxx
	PLACE1:		
20	PLACE1xxxxxx	F-PLACE1xxxxxx	R-PLACE1xxxxxx
	PLACE2:		
	PLACE2xxxxxx	F-PLACE2xxxxxx	R-PLACE2xxxxx
•	PLACE3:		
25	PLACE3xxxxxx	F-PLACE3xxxxxx	R-PLACE3xxxxxx
	PLACE4:		D DI 1051
	PLACE4xxxxxx	F-PLACE4xxxxxx	R-PLACE4xxxxxx
30	OVARC1:	E 014001	D OVAROLISM
30	OVARC1xxxxxx	F-0VARC1xxxxxx	R-OVARC1xxxxxx
	HEMBA1:	F-HEMBA1xxxxxx	R-HEMBA1xxxxxx
	HEMBA1xxxxxx HEMBB1:	r-nembalxxxxxx	K-UCINDA I XXXXXX
35	HEMBB1xxxxxx	F-HEMBB1xxxxxx	R-HEMBB1xxxxxx
	MAMMA1:	I IILIIDDIXXXXX	II IILMDDIXXXXX
	MAMMA1xxxxx	F-MAMMA1xxxxxx	R-MAMMA1xxxxxx
	THYRO1:	1 HB UMB T AXXXXX	
40	THYRO1xxxxxx	F-THYR01xxxxxx	R-THYR01xxxxxx
	VESEN1:		
	VESEN1xxxxxx	F-VESEN1xxxxxx	R-VESEN1xxxxxx
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# **EXAMPLE 2**

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Estimation of the fullness ratio at the 5'-ends of the clones contained in the cDNA libraries constructed by the oligocapping method.

[0139] The fullness ratio at the 5'-end sequence of the 59,823 clones in the human cDNA libraries constructed by the oligo-capping method was determined as follows. Of all the clones whose 5'-end sequences were found in those of known human mRNA in the public database, a clone was judged to be "full-length", if it had a longer 5'-end sequence than that of the known human mRNA, or, even though the 5'-end sequence was shorter, if it contained the translation initiation codon. A clone which did not contain the translation initiation codon was judged to be "not-full-length". The fullness ratio ((the number of full-length clones)/(the number of full-length and not-full-length clones)) at the 5'-end of the cDNA clones from each library was determined by comparing with the known human mRNA. As a result, the fullness

ratio of the 5'-ends was 63.5%. The result indicates that the fullness ratio at the 5'-end sequence was extremely high.

# **EXAMPLE 3**

Assessment of the fullness ratio of the 5'-end of the cDNA by the ATGpr and the ESTiMateFL.

[0140] The ATGpr, developed by Salamov A.A., Nishikawa T., and Swindells M.B. in the Helix Research Institute, is a program for prediction of the translation initiation codon based on the characteristics of the sequences in the vicinity of the ATG codon. The results are shown with expectations (also described as ATGpr1 below) that an ATG is a true initiation codon (0.05-0.94) (can be described as ATGpr1). When the program was applied to the 5'-end sequences of the clones from the cDNA library that was obtained by the oligo-capping method and that had 65% fullness ratio, the sensitivity and specificity of estimation of a full-length clone (clone containing the N-terminal end of ORF) were improved to 82-83% by selecting only clones having the ATGpr1 score 0.6 or higher.

[0141] Furthermore, the program was used to assess the fullness of 18,959 clones in the human libraries obtained here, which have 5'-ends matched to a known human mRNA.

[0142] Briefly, the maximal ATGpr1 score of the clones was determined, and then their 5'-end sequence was compared with the known human mRNA to estimate whether the clone is full-length or not. The result is shown in Table 4. [0143] Based on the knowledge that known mRNAs, in general, are highly expressed in the cell, similar estimation was performed with genes having a low number in the EST hit, which represent relatively low abundant mRNAs, and the result is shown in Table 5.

[0144] In the table, the number of full-length clones indicate that of clones containing the N-terminal end of ORF, and so does the number of not-full-length clones that of clones without the N-terminal end of ORF. The fullness ratio represents (the number of full-length clones)/(the number of full-length clones plus the number of not-full-length clones).

Table 4

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having a matched 5'-end with that of a known mRNA.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	11,193	9,346	83.5%
>=0.50	13,369	10,549	78.9%
>=0.30	15,489	11,340	73.2%
>=0.15	17,394	11,811	67.9%
>=0.00	18,959	12,046	63.5%

Table 5

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of the clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having 5 EST hits or less among the clones having a matched 5'-end with that of a known mRNA.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	2,801	1,934	69.0%
>=0.50	3,683	2,393	65.0%
>=0.30	4,683	2,707	57.8%
>=0.15	5,559	2,890	52.0%
>=0.00	6,113	3,013	49.8%

[0145] Next, the ESTiMateFL was used for the estimation. The ESTiMateFL, developed by Nishikawa and Ota in the Helix Research Institute, is a method for the selection of a clone with high fullness ratio by comparing with the 5'-end or 3'-end sequences of ESTs in the public database.

[0146] By the method, a cDNA clone is judged to be most likely not to be full-length if there exist any ESTs which

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have longer 5'-end or 3'-end sequences than the clone. The method is systematized for high throughput analysis. A clone is judged to be full-length if the clone has a longer 5'-end sequence than ESTs in the public database. Even if a clone has a shorter 5'-end, the clone is judged to be full-length if the difference in length is within 50 bases, and otherwise judged not to be full-length, for convenience.

[0147] In case of the clones whose 5'-end sequence is matching with the known mRNA, 80% of the clones judged to be full-length by comparing with ESTs was also judged to be full-length by comparing with the known mRNA. Also, 80% of the clones judged to be not full-length by comparing with ESTs was also judged to be not full-length by comparing with the known mRNA.

[0148] The precision of the estimation by comparing with ESTs is improved with increasing number of ESTs to be compared. However, in case that a limited number of ESTs are available, the reliability becomes low. Thus, the method is effective in excluding clones with high probability of being not-full-length, from the cDNA clones that is synthesized by the oligo-capping method and that have the 5'-end sequences with about 60 % fullness ratio. In particular, the ESTiMateFL is efficiently used to estimate the fullness ratio at the 3'-end sequence of cDNA of a human unknown mRNA which has a significant number of EST deposits in the public database.

[0149] The 18,959 clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, were estimated by using the ATGpr and ESTiMateFL. Briefly, the 5'-end sequence of the respective clone was analyzed to obtain the maximal ATGpr1 score, and compared with the ORF of the known human mRNA that matches with it to determine whether the clone is full-length or not. Then, the 5'-end sequence of the respective clone was analyzed by the ESTiMateFL to judge whether the clone is full-length or not. Specifically, the 5'-end sequences of the 18,959 clones were compared with those of ESTs by the ESTiMateFL and the clones other than those that are not full-length were selected. Then, the selected clones were used to analyze the relationship between the ATGpr and the fullness ratio. The result was summarized in Table 6. Also, among the selected, the clones in which the number of the EST hit is not more than 5 were selected and analyzed. The result was summarized in Table 7, which represents the result of the analysis of mRNA with relatively low abundance.

[0150] Therein, the number of being full-length, the number of being not full-length, and the fullness ratio indicate the number of the clones that contain the N-terminus of the ORF, the number of the clones that do not contain the N-terminus of the ORF, and (the number of being full-length)/(the number of being full-length) plus (the number of being not full-length), respectively.

Table 6

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequence in the clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, and also other than those being not full-length according to the comparison with ESTs.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	9,068	8,349	92.1%
>=0.50	10,345	9,318	90.1%
>=0.30	11,425	9,964	87.2%
>=0.15	12,254	10,335	84.3%
>=0.00	12,785	10,484	82.0%

Table 7

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	1,959	1,510	77.1%
>=0.50	2,469	1,821	73.8%
>=0.30	2,975	2,046	68.8%
>=0.15	3,368	2,164	64.3%

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# Table 7 (continued)

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.00	3,661	2,226	60.8%

[0151] The 19,226 clones, isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to that of a known human mRNA were estimated by the ATGpr2, and the correlation between the score and the fullness ratio was estimated. Specifically, the maximal ATGpr2 score of the clones identical to a known human mRNA was determined, and then their fullness ratio was estimated by comparing the 5'-ends with ORF of known human mRNA. The result was shown in Table 8.

## Table 8

Maximal ATGpr2 score and fullness ratio of the 5'-end sequence of the clones, which are isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA.

maximal ATGpr2 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.30	10,748	8,031	74.7%
>=0.15	16,383	11,226	68.5%
>=0.00	19,226	12,285	63.9%

[0152] According to the above results, it was found that, in case of using clones isolated from human cDNA libraries constructed by the oligo-capping method, the fullness ratio of the clones that have low score in the ATGpr can be improved by estimating their 5'-end sequence using the combination of the ATGpr and the ESTiMateFL. Therefore, the method was applied to select a cDNA clone with high fullness ratio.

# 35 EXAMPLE 4

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Clustering of the 5'-end and 3'-end sequences of cDNA clones.

[0153] The 5'-end and 3'-end sequences of cDNA clones were obtained, and clustered separately. The single pass data of the nucleotide sequence of the 5'-end and 3'-end was subject to the BLAST search between the sequence data of all the clones synthesized in example 1, and the clones considered to be originating from the same gene were put together into a group. If the 5'-end of a clone contains the consensus sequence of 300 bases or more with identity 95% or more, or the 3'-end contains the consensus sequence of 200 bases or more and having identity 90% or more, the clones were put in the same group.

<sup>45</sup> [0154] The groups of the 5'-end sequence and the 3'-end sequence were further clustered so as that the groups from the same clone can be in the same group (cluster).

# **EXAMPLE 5**

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50 Characterization of the cloned sequence.

[0155] The data of the 5'-end sequence of the cloned sequence was characterized by the following way:

- (1) examining whether it is identical to those of mRNA from human and other species (including authorized sequences) and human EST by the BLAST homology search of the GenBank,
- (2) examining whether it has longer 5'-end than those of human mRNA and human EST,
- (3) determining the scores in the ATGpr1 and ATGpr2 programs of all the initiation codons in the 5'-end sequence, and

- (4) determining the number of the human EST clone(s) that is judged to be identical by the BLAST homology search of the GenBank.
- [0156] The data of the 3'-end sequence of the cloned sequence was characterized by the above (1) and (4).
- [0157] These characterized data were used for the final selection of the clones.

# **EXAMPLE 6**

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Identity to the human mRNA and human EST, and comparison of the length of the 5'-end.

[0158] The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of mRNA from human or other species when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 94% or more. The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of human EST when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 90% or more.

[0159] The sequence of the clone was judged to be full-length in comparison with human mRNA when the sequence has longer 5'-end, or it contains the translation initiation site. The sequence of the clone was judged to be full-length in comparison with human EST in the database when the sequence has longer 5'-end, or while it has shorter end, the difference in length between the two sequences is 50 bases or less. The other clones were judged to be not full-length.

# EXAMPLE 7

Prediction of the fullness ratio by the ATGpr.

25 [0160] The score in the ATGpr1 is the expectation to be full-length based on calculations, and the higher score reflects the higher probability to be full-length as shown in Example 3. The maximal ATGpr1 score and the maximal ATGpr2 score represent the score obtained with all the initiation codons contained in the 5'-end sequence of the cloned sequence, and were used for the characterization.

# 30 EXAMPLE 8

Prediction of the novelty using the number of the identical ESTs by the homology search.

[0161] For both the 5'-end and 3'-end sequences of the clones, the number of the identical ESTs was determined by the homology search on the GenBank. Human ESTs were judged to be identical when the EST has a sequence of 200 nucleotides or more with 90% or more matching with the 5'-end sequence. The number of the identical ESTs were used for characterization and as an index of novelty. The clone having not identical sequence at the 5'-end and 3'-end sequences to those of mRNA as well as those of ESTs is a gene encoding a novel protein. Similarly, a clone having either the 5'-end or the 3'-end sequences, which has low number of the identical ESTs, is judged to be a gene encoding a novel protein.

# **EXAMPLE 9**

Characterization of clusters.

[0162] The clusters of the groups of the 5'-end and 3'-end sequences were characterized according to the following criteria.

(1) Whether it is identical to the mRNA sequences from human or other species (including authorized sequences), or human ESTs by the BLAST search of the GenBank.

A cluster containing at least one sequence of all the 5'-end and 3'-end sequences, which is identical to one of the mRNA sequences, was regarded to be the same cluster of the mRNA sequence.

(2) Whether it has longer 5'-end than human mRNA sequence and human ESTs.

When all the 5'-end sequences contained in a cluster are judged to be not full-length compared with the mRNA sequences and human ESTs, the cluster was regarded as being not full-length.

(3) The scores in the ATGpr1 and ATGpr2 using all the initiation codons contained in the 5'-end sequences. The maximal ATGpr1 score among those of all the 5'-end sequences in a cluster was determined as the ATGpr1 score of the cluster. The ATGpr2 score of the cluster was also determined in the same way.

- (4) The number of the identical human ESTs determined by the BLAST search of the GenBank.
- [0163] The maximum number was determined in the numbers of ESTs identical to each of 5'-end sequences contained in a cluster. The number of the ESTs identical to the 5'-end sequences in the cluster was defined as the maximum number. The number of the ESTs identical to the 3'-end sequences in a cluster was determined in the same way.

## **EXAMPLE 10**

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Methods for selection of the clusters by the characteristics.

[0164] Data obtained by the characterization described above was used to discard the clusters that are identical to any mRNA sequence from human and other species (including authorized sequences), or those clusters that are not full-length. From the rest of the clusters, the clusters that fulfill any of the following conditions were selected.

- (a) A cluster in which the number of the identical ESTs for the 5'-end sequence is 20 or less, and the ATGpr1 score exceeds 0.3.
- (b) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for both the 5'-end sequence and the 3'-end sequence is 5 or less, and multiple clones are contained.
- (c) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is 0, and the number of the identical ESTs for the 3'-end sequence is not less than 1.
- (d) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is not less than 1 and not more than 5, and the number of the identical ESTs for the 3'-end sequence is 0.
- [0165] The clusters selected by (a) contain at least one clone that is novel and having high fullness ratio. The clusters selected by (b), (c), and (d) contain at least one clone that is novel and having low fullness ratio, but is still full-length.

## **EXAMPLE 11**

Methods for selection of clones from clusters.

[0166] In the clusters comprising a single clone, the clone was selected.

[0167] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score higher than 0.3, a clone with the highest score was selected.

[0168] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score not more than 0.3, a clone with the highest ATGpr2 score was selected, if the score was higher than 0.3.

[0169] In the clusters comprising multiple clones, in which the clones have the scores not more than 0.3 in both the ATGpr1 and the ATGpr2, a clone with the highest scores in both the ATGpr1 and ATGpr2 was selected.

[0170] In the clusters comprising multiple clones, in which the above selection by the ATGpr score was not applicable, selected was a clone having longer 5'-end by assembling the 5'-end sequence, 3'-end sequence, and human ESTs. For assembling, the Sequencher™ (Hitachi Soft Engineering) was used. When even the selection by assembling failed, all the clones were judged to be full-length.

[0171] As a result, 3690 clones were the clones that have the maximal ATGpr1 score higher than 0.3. On the other hand, 477 clones were the clones that have the maximal ATGpr1 score not more than 0.3, and the maximal ATGpr2 score higher than 0.3. The number of the clones having the highest scores in both the ATGpr1 and ATGpr2, while the scores were not more than 0.3, were 97. The number of the clones which were not selected by the ATGpr scores, but were selected by assembling the 5'-end sequence, 3'-end sequence, and human ESTs, were 117. The clones that have the score in both the ATGpr1 and ATGpr2 not more than 0.3, but were selected because the cluster comprises a single clone, were 1166. In the clones, at least either of the 5'-end or 3'-end sequence was not identical to any of human ESTs. Some clones were selected because the cluster comprises a single clone, or by assembling, in which there is no ATG codon (9 clones: HEMBA1001960, HEMBA106569, HEMBB1001454, NT2PR2002839, NT2RP2005325, NT2RP2006323, PLACE1004506, PLACE1005526, and THYRO1001177). The sequences that do not contain the ATG codon were considered to be corresponding to the 5'-UTR. Although the clones do not have the scores in the ATGpr1 and ATGpr2, the clones were yet judged to be full-length according to the fullness ratio, as shown in Table 4, 5, 6, 7, and 8. The above clones that were finally judged to be full-length were classified into 11 groups according to the following criteria.

Group (1): 1516 clones

Among the 3690 clones having the maximal ATGpr1 score higher than 0.3, the following 1516 clones were having

high fullness ratio and a novel clone, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs. HEMBA1000046, HEMBA1000050, HEMBA1000129, HEMBA1000150, HEMBA1000158, HEMBA1000193 , HEMBA1000201, HEMBA1000216, HEMBA1000227, HEMBA1000288, HEMBA1000290, HEMBA100030 3, HEMBA1000304, HEMBA1000369, HEMBA1000392, HEMBA1000396, HEMBA1000488, HEMBA10005 05, HEMBA1000508, HEMBA1000534, HEMBA1000542, HEMBA1000594, HEMBA1000637, HEMBA1000 657, HEMBA1000752, HEMBA1000867, HEMBA1000869, HEMBA1000872, HEMBA1000910, HEMBA100 0918, HEMBA1000919, HEMBA1000942, HEMBA1000968, HEMBA1000975, HEMBA1000986, HEMBA10 01022, HEMBA1001043, HEMBA1001052, HEMBA1001080, HEMBA1001085, HEMBA1001109, HEMBA1 001140, HEMBA1001174, HEMBA1001235, HEMBA1001286, HEMBA1001302, HEMBA1001398, HEMBA 1001407, HEMBA1001415, HEMBA1001446, HEMBA1001476, HEMBA1001497, HEMBA1001510, HEMB A1001533, HEMBA1001570, HEMBA1001581, HEMBA1001635, HEMBA1001640, HEMBA1001647, HEM BA1001661, HEMBA1001731, HEMBA1001744, HEMBA1001746, HEMBA1001800, HEMBA1001815, HE MBA1001822, HEMBA1001866, HEMBA1001896, HEMBA1001910, HEMBA1001987. HEMBA1002018, HEMBA1002035, HEMBA1002049, HEMBA1002092, HEMBA1002119, HEMBA1002125, HEMBA1002161, HEMBA1002177, HEMBA1002189, HEMBA1002191, HEMBA1002199, HEMBA100222 9, HEMBA1002237, HEMBA1002265, HEMBA1002363, HEMBA1002417, HEMBA1002419, HEMBA10024 30, HEMBA1002439, HEMBA1002477, HEMBA1002503, HEMBA1002508, HEMBA1002515, HEMBA1002 547, HEMBA1002688, HEMBA1002703, HEMBA1002746, HEMBA1002750, HEMBA1002850, HEMBA100 2973, HEMBA1003021, HEMBA1003067, HEMBA1003077, HEMBA1003078, HEMBA1003079, HEMBA10 03117, HEMBA1003129, HEMBA1003175, HEMBA1003199, HEMBA1003235, HEMBA1003250, HEMBA1 003257, HEMBA1003291, HEMBA1003322, HEMBA1003327, HEMBA1003370, HEMBA1003380, HEMBA 1003395, HEMBA1003402, HEMBA1003461, HEMBA1003480, HEMBA1003538, HEMBA1003545, HEMB A1003556, HEMBA1003581, HEMBA1003621, HEMBA1003645, HEMBA1003667, HEMBA1003720, HEM BA1003760, HEMBA1003799, HEMBA1003807, HEMBA1003827, HEMBA1003836, HEMBA1003866, HE MBA1003879, HEMBA1003880, HEMBA1003985, HEMBA1003989, HEMBA1004011, HEMBA1004048, HEMBA1004056, HEMBA1004074, HEMBA1004133, HEMBA1004146, HEMBA1004150, HEMBA1004199, HEMBA1004200, HEMBA1004238, HEMBA1004246, HEMBA100427 5, HEMBA1004286, HEMBA1004289, HEMBA1004321, HEMBA1004327, HEMBA1004335, HEMBA10043 41, HEMBA1004372, HEMBA1004389, HEMBA1004479, HEMBA1004499, HEMBA1004507, HEMBA1004 542, HEMBA1004554, HEMBA1004573, HEMBA1004596, HEMBA1004632, HEMBA1004697, HEMBA100 4705, HEMBA1004709, HEMBA1004711, HEMBA1004736, HEMBA1004751, HEMBA1004752, HEMBA10 04753, HEMBA1004756, HEMBA1004758, HEMBA1004768, HEMBA1004771, HEMBA1004795, HEMBA1 004806, HEMBA1004850, HEMBA1004863, HEMBA1004889, HEMBA1004923, HEMBA1004929, HEMBA 1004930, HEMBA1004944, HEMBA1004972, HEMBA1004980, HEMBA1005019, HEMBA1005035, HEMB A1005050, HEMBA1005066, HEMBA1005075, HEMBA1005079, HEMBA1005083, HEMBA1005113, HEM BA1005133, HEMBA1005149, HEMBA1005219, HEMBA1005331, HEMBA1005338, HEMBA1005367, HE MBA1005411, HEMBA1005468, HEMBA1005469, HEMBA1005517, HEMBA1005518, HEMBA1005526, H EMBA1005548, HEMBA1005576, HEMBA1005583, HEMBA1005595, HEMBA1005609, HEMBA1005685, HEMBA1005732, HEMBA1005755, HEMBA1005813, HEMBA1005815, HEMBA1005834, HEMBA1005884, HEMBA1005963, HEMBA1005991, HEMBA1006031, HEMBA1006100, HEMBA1006121, HEMBA1006138, HEMBA1006173, HEMBA1006182, HEMBA1006198, HEMBA1006252, HEMBA1006272, HEMBA1006278, HEMBA1006291, HEMBA100629 3, HEMBA1006344, HEMBA1006349, HEMBA1006377, HEMBA1006381, HEMBA1006398, HEMBA10064 24, HEMBA1006467, HEMBA1006474, HEMBA1006483, HEMBA1006492, HEMBA1006494, HEMBA1006 497, HEMBA1006502, HEMBA1006530, HEMBA1006579, HEMBA1006583, HEMBA1006643, HEMBA100 6674, HEMBA1006682, HEMBA1006709, HEMBA1006717, HEMBA1006737, HEMBA1006754, HEMBA10 06758, HEMBA1006767, HEMBA1006795, HEMBA1006796, HEMBA1006807, HEMBA1006832, HEMBA1 006900, HEMBA1006973, HEMBA1006976, HEMBA1006993, HEMBA1006996, HEMBA1007002, HEMBA 1007052, HEMBA1007062, HEMBA1007066, HEMBA1007151, HEMBA1007203, HEMBA1007281, HEMB A1007300, HEMBA1007320, HEMBA1007342, HEMBB1000008, HEMBB1000024, HEMBB1000025, HEMBB1000083, HEMBB1000103, HEMBB1000173, HEMBB1000175, HEMBB1000198, HEMBB1000240, HEMBB1000244, HEMBB1000338, HEMBB100033 9, HEMBB1000391, HEMBB1000438, HEMBB1000510, HEMBB1000550, HEMBB1000556, HEMBB10005 89, HEMBB1000591, HEMBB1000593, HEMBB1000632, HEMBB1000671, HEMBB1000673, HEMBB1000 693, HEMBB1000706, HEMBB1000725, HEMBB1000781, HEMBB1000810, HEMBB1000826, HEMBB100 0835, HEMBB1000848, HEMBB1000852, HEMBB1000870, HEMBB1000887, HEMBB1000908, HEMBB10 00927, HEMBB1000947, HEMBB1000973, HEMBB1000991, HEMBB1001011, HEMBB1001014, HEMBB1 001020,

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## Group (2): 377 clones

35

Among the 3690 clones, the following 377 clones were full-length, and a novel clone, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

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## 50 Group (3): 1797 clones

Among the 3690 clones, the following 1797 clones were full-length, and novel clones, in which the number of the identical human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

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PLACE4000089, PLACE4000106, PLACE4000259, PLACE4000269, PLACE4000326, PLACE400 0369,
PLACE4000431, PLACE4000445, PLACE4000581, PLACE4000593, PLACE4000650, PLACE40 00670,
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THYRO1000173, THYRO1000197, THYRO1000199, THYRO1000206, THYRO1000242, THYRO100027 0,
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THYRO1000488, THYRO1000502, THYRO1000569, THYRO1000570, THYRO1000585, THYRO1000 605,
THYRO1000715, THYRO1000756, THYRO1000777, THYRO1000829, THYRO1000855, THYRO100 0926,
THYRO1000983, THYRO1000984, THYRO1001120, THYRO1001134, THYRO1001173, THYRO10 01204,
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THYRO1001793,
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                               Y79AA1000181,
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Y79AA1000231,
              Y79AA1000258,
                              Y79AA1000313,
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Y79AA1000574,
              Y79AA1000589,
                              Y79AA1000705.
                                             Y79AA1000734,
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                                                                            Y79AA100 0833.
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                              Y79AA1001023,
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Y79AA1002089.
              Y79AA1002093.
                              Y79AA1002125.
                                             Y79AA1002208,
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Y79AA1002234.
              Y79AA1002258.
                              Y79AA1002307,
                                             Y79AA1002311,
                                                            Y79AA1002351,
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Y79AA1002487.
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### Group (4): 453 clones

20

Among the 1857 clones having the maximal ATGpr1 score not more than 0.3 (including the 9 clones whose 5'-end sequence does not contain the ATG codon), the following 453 clones were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

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HEMBA1000180, HEMBA1000280, HEMBA1000282, HEMBA1000333, HEMBA1000351, HEMBA1000357,
HEMBA1000376, HEMBA1000469, HEMBA1000504, HEMBA1000519, HEMBA1000545, HEMBA100055 7,
HEMBA1000575, HEMBA1000726, HEMBA1000822, HEMBA1000934, HEMBA1000943, HEMBA10009 60,
HEMBA1000972, HEMBA1001020, HEMBA1001051, HEMBA1001060, HEMBA1001071, HEMBA1001 208,
HEMBA1001319, HEMBA1001383, HEMBA1001411, HEMBA1001433, HEMBA1001435, HEMBA100 1478,
HEMBA1001522, HEMBA1001636, HEMBA1001651, HEMBA1001658, HEMBA1001709, HEMBA10 01745,
HEMBA1001791, HEMBA1001835, HEMBA1001844, HEMBA1001888, HEMBA1001940, HEMBA1 001962,
HEMBA1002022, HEMBA1002185, HEMBA1002348, HEMBA1002621, HEMBA1002645, HEMBA 1002661,
HEMBA1002728, HEMBA1002921, HEMBA1002924, HEMBA1002934, HEMBA1002968, HEMBA1003037,
HEMBA1003071, HEMBA1003166, HEMBA1003202, HEMBA1003204, HEMBA1003220, HEMBA1003229,
HEMBA1003276, HEMBA1003296, HEMBA1003373, HEMBA1003531, HEMBA100364 0, HEMBA1003856,
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HEMBA1004433, HEMBA1004577, HEMBA1004730, HEMBA1004778, HEMBA1004 803, HEMBA1004807,
HEMBA1004880, HEMBA1004900, HEMBA1004983, HEMBA1005123, HEMBA100 5241, HEMBA1005311,
HEMBA1005318, HEMBA1005353, HEMBA1005374, HEMBA1005447, HEMBA10 05588, HEMBA1005593,
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HEMBA1006253, HEMBA1006259, HEMBA1006364, HEMBA1006380, HEMBA 1006426, HEMBA1006562,
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HEMBA1006949, HEMBA1007078, HEMBA1007129, HEMBA1007147, HEM BA1007206, HEMBA1007279,
HEMBA1007327,
HEMBB1000005, HEMBB1000055, HEMBB1000144, HEMBB1000258, HEMBB1000318, HEMBB1000335,
HEMBB1000354, HEMBB1000374, HEMBB1000402, HEMBB1000404, HEMBB1000480, HEMBB100049 3,
HEMBB1000554. HEMBB1000573, HEMBB1000649, HEMBB1000652, HEMBB1000709, HEMBB10007 49,
HEMBB1000790, HEMBB1000827, HEMBB1000831, HEMBB1000893, HEMBB1001004, HEMBB1001 008,
HEMBB1001047, HEMBB1001315, HEMBB1001317, HEMBB1001326, HEMBB1001367, HEMBB100 1424,
HEMBB1001436, HEMBB1001458, HEMBB1001535, HEMBB1001565, HEMBB1001747, HEMBB10 01749,
HEMBB1001797, HEMBB1001836, HEMBB1001863, HEMBB1001875, HEMBB1001911, HEMBB1 001922,
HEMBB1001925, HEMBB1001944, HEMBB1001983, HEMBB1001996, HEMBB1001997, HEMBB 1002092,
HEMBB1002247, HEMBB1002266, HEMBB1002387, HEMBB1002425, HEMBB1002458, HEMB B1002522.
HEMBB1002534, HEMBB1002582, HEMBB1002596, HEMBB1002617, HEMBB1002702, MAMMA1000043,
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MAMMA1000360, MAMMA1000402, MAMMA1000414, MAMMA1000500, MAMMA100052 2, MAMMA1000576,
MAMMA1000594, MAMMA1000597, MAMMA1000720, MAMMA1000775, MAMMA10007 78, MAMMA1000798,
MAMMA1000862. MAMMA1000876, MAMMA1000931, MAMMA1000940, MAMMA1000 941, MAMMA1000975,
MAMMA1001038, MAMMA1001186, MAMMA1001220, MAMMA1001256, MAMMA100 1274, MAMMA1001341,
MAMMA1001397, MAMMA1001420, MAMMA1001547, MAMMA1001670, MAMMA10 01679, MAMMA1001711,
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MAMMA1001745, MAMMA1001760, MAMMA1001769, MAMMA1001815, MAMMA1 001907, MAMMA1002056,
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MAMMA1002311, MAMMA1002411, MAMMA1002498, MAMMA1002571, MAMM A1002701, MAMMA1002727,
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MAMMA1002909, MAMMA1002941, MAMMA1002973, MAMMA1003004, MA MMA1003007, MAMMA1003039,
MAMMA1003089.
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NT2RP2003000,
              NT2RP2003073,
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                                           PLACE3000271, PLACE3000373, PLACE3000399.
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THYRO1000865, THYRO1001003, THYRO1001031, THYRO1001133, THYRO1001401, THYRO100142 6,
THYRO1001434, THYRO1001559, THYRO1001570, THYRO1001706, THYRO1001746, THYRO10017 72,
THYRO1001907, Y79AA1000033, Y79AA1000346, Y79AA1000805, Y79AA1001692, Y79AA1002 220.
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## Group (5): 24 clones

45

The following 24 clones having low score in the ATGpr1 were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000622, HEMBA1000749, HEMBA1000876, HEMBA1001226, HEMBA1001391, HEMBA1002742, HEMBA1003908, HEMBA1004000, HEMBB1000336, MAMMA1000356, MAMMA1000883, MAMMA100159 0, NT2RP2000289, NT2RP2001467, PLACE1002853, PLACE1003420, PLACE1004836, PLACE10049 13, PLACE1006795, PLACE1008181, PLACE1008715, PLACE4000344, THYRO1000129, THYRO1001 321.

## Group (6): 65 clones

The following 65 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

HEMBA1000604, HEMBA1000673, HEMBA1001024, HEMBA1001026, HEMBA1001734, HEMBA1001784,

HEMBA1001808, HEMBA1003902, HEMBA1004164, HEMBA1004909, HEMBA1005232, HEMBA100557 7, HEMBA1006461, HEMBA1006695, HEMBB10011119, HEMBB1001337, HEMBB1001536, HEMBB100118 68, HEMBB1002045, HEMBB1002579, MAMMA1000444, MAMMA1000761, MAMMA1000943, MAMMA1001 820, MAMMA1002360,

NT2RM4000366, NT2RM4001856, NT2RM4002390, NT2RP2000108, NT2RP2000257, NT2RP2001506, NT2RP2002047, NT2RP200266, NT2RP2002475, NT2RP2004400, NT2RP2004587, NT2RP200528 9, NT2RP2005694, NT2RP3001898, NT2RP3003264, NT2RP3003433, NT2RP3003842, OVARC10012 40, PLACE1001323, PLACE1002227, PLACE1002500, PLACE1002604, PLACE1002772, PLACE1003 478, PLACE1004681, PLACE1005108, PLACE1005932, PLACE1006318, PLACE1006368, PLACE100 6506, PLACE1006904, PLACE1007557, PLACE1007877, PLACE1009048, PLACE1011109, PLACE10 11643, PLACE4000548, THYRO1000279, Y79AA1000410, Y79AA1002103,

## Group (7): 32 clones

The following 32 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA100251, HEMBA1001803, HEMBA1001918, HEMBA1002257, HEMBA1003064, HEMBA1003714, HEMBA1004405, HEMBA1005508, HEMBB1000054, HEMBB1001142, MAMMA1000175, MAMMA100116 2, MAMMA1002972, NT2RM4000425, NT2RP2004512, NT2RP2005531, NT2RP2005942, NT2RP20065 54, NT2RP3001007, NT2RP3001318, OVARC1000017, OVARC1000068, OVARC1000486, PLACE1001 705, PLACE1002319, PLACE1007743, PLACE1007829, PLACE1008630, PLACE1009925, PLACE101 1492, PLACE1011749, THYRO1000793,

The following 117 clones, selected by assembling the sequence of the other clones in the same cluster and human ESTs, have high fullness ratio. The clones were classified into the following group (8) and (9). HEMBA1001323, HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1002538, HEMBA1003309, HEMBA1003939, HEMBA1004015, HEMBA1004295, HEMBA100467 2, HEMBA1004865, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1006779, HEMBA10072 88, HEMBB1000218, HEMBB1000272, HEMBB1000399, HEMBB1000491, HEMBB1000996, HEMBB1001 114, HEMBB1001850, HEMBB1002015, MAMMA1000287, MAMMA1001683, MAMMA1001686, MAMMA100 2612, NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP2000270, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002736, NT2RP2002753, NT2RP2003456, NT2RP200372 7, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2004580, NT2RP20052 93, NT2RP2005476, NT2RP2005753, NT2RP2005815, NT2RP2005841, NT2RP2005857, NT2RP2006 393, NT2RP2006467, NT2RP3000109, NT2RP3000449, NT2RP3001245, NT2RP3001634, NT2RP300 2056, NT2RP3002810, NT2RP3002955, NT2RP3003032, NT2RP3003138, NT2RP3003500, NT2RP30 03819, NT2RP4000078, NT2RP4000515, NT2RP4000517, NT2RP4001407, NT2RP4001889, NT2RP4 002905, OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691, PLACE1001746, PLACE1001748, PLACE1001845, PLACE1002066, PLACE1003373, PLACE100390 0, PLACE1004118, PLACE1004256, PLACE1004284, PLACE1004336, PLACE1004506, PLACE10049 34, PLACE1005077, PLACE1005111, PLACE1005409, PLACE1005730, PLACE1006076, PLACE1006 360, PLACE1006470, PLACE1006760, PLACE1006867, PLACE1007045, PLACE1007111, PLACE100 7807, PLACE1008080, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008426, PLACE10 08621, PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, PLACE3000276, THYRO1 000805,

## Group (8): 36 clones

THYRO1001365, THYRO1001673, Y79AA1001848,

Among the 117 clones described above, the following 36 clones were novel clones, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs.

HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1003939, HEMBA1004295, HEMBA1004865, HEMBA1006779, HEMBB1000218, HEMBB1000491, HEMBB100099 6, HEMBB1001114, MAMMA1000287, MAMMA1001686, MAMMA1002612, NT2RP2002736, NT2RP20037 27, NT2RP2004580, NT2RP2005841, NT2RP2006393, PLACE1002066, PLACE1003373, PLACE1003 900, PLACE1004118, PLACE1004336, PLACE1004506, PLACE1004934, PLACE1005409, PLACE100 5730, PLACE1006470, PLACE1008080, PLACE3000276, THYRO1000805, THYRO1001365, THYRO10 01673,

## Group (9): 81 clones

Among the 117 clones described above, the following 81 clones were the clones in which the number of the identical

human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

HEMBA1001323, HEMBA1002538, HEMBA1003309, HEMBA1004015, HEMBA1004672, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1007288, HEMBB1000272, HEMBB1000399, HEMBB100185 0, HEMBB1002015, MAMMA1001683, NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP20002 70, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002753, NT2RP2003 456. NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2005293, NT2RP200 5476. NT2RP2005753, NT2RP2005815, NT2RP2005857, NT2RP2006467, NT2RP3000109, NT2RP30 00449. NT2RP3001245, NT2RP3001634, NT2RP3002056, NT2RP3002810, NT2RP3002955, NT2RP3 003032. NT2RP3003138, NT2RP3003500, NT2RP3003819, NT2RP4000078, NT2RP4000515, NT2RP 4000517, NT2RP4001407, NT2RP4001889, NT2RP4002905, OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691, PLACE1001746, PLACE1001748, PLACE1001845, PLACE1004256, PLACE1004284, PLACE100507 7, PLACE1005111, PLACE1006076, PLACE1006360, PLACE1006760, PLACE1006867, PLACE10070 45,

### Group (10): 938 clones

20

The following 938 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 4. The clones were novel clones, in which at least the 5'-end sequence was not identical to any of human ESTs.

PLACE1007111, PLACE1007807, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008 426,

PLACE1008621, PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, Y79AA100 1848,

HEMBA100042, HEMBA1000213, HEMBA1000264, HEMBA1000355, HEMBA1000366, HEMBA1000411, HEMBA1000422, HEMBA1000428, HEMBA1000434, HEMBA1000442, HEMBA1000464, HEMBA100054 0, HEMBA1000763, HEMBA1000747, HEMBA1000655, HEMBA1000662, HEMBA1000702, HEMBA10007 05, HEMBA1000843, HEMBA1000852, HEMBA1000870, HEMBA1000908, HEMBA1000908, HEMBA1001017, HEMBA1001019, HEMBA1001094, HEMBA1001121, HEMBA1001172, HEMBA1001375, HEMBA1001432, HEMBA1001454, HEMBA1001463, HEMBA1001515, HEMBA1001557, HEMBA1001566, HEMBA1001585, HEMBA1001589, HEMBA1001678, HEMBA1001678, HEMBA1001681, HEMBA1001718, HEMBA1001761, HEMBA1001945, HEMBA1001960, HEMBA1001964, HEMBA1001991,

HEMBA1002003, HEMBA1002008, HEMBA1002039, HEMBA1002100, HEMBA1002113, HEMBA1002139, HEMBA1002144, HEMBA1002153, HEMBA1002226, HEMBA1002253, HEMBA1002270, HEMBA100232 1, HEMBA1002328, HEMBA1002337, HEMBA1002381, HEMBA1002486, HEMBA1002552, HEMBA1002558, HEMBA1002561, HEMBA1002590, HEMBA1002628, HEMBA1002629, HEMBA1002659, HEMBA1002666, HEMBA1002678, HEMBA1002712, HEMBA1002716, HEMBA1002730, HEMBA1002748, HEMBA1002780, HEMBA1003801, HEMBA1003803, HEMBA1003083, HEMBA1003083, HEMBA1003083, HEMBA1003084, HEMBA1003098, HEMBA1003314, HEMBA1003273, HEMBA1003373, HEMBA1003376, HEMBA1003334, HEMBA1003348, HEMBA1003376, HEMBA1003758, HEMBA1003773, HEMBA1003784, HEMBA1003784, HEMBA1003937, HEMBA1003937, HEMBA1003773, HEMBA1003784, HEMBA1003885, HEMBA1003937, HEMBA1003950,

HEMBA1004012, HEMBA1004045, HEMBA1004138, HEMBA1004241, HEMBA1004272, HEMBA1004306, HEMBA1004312, HEMBA1004334, HEMBA1004354, HEMBA1004356, HEMBA1004366, HEMBA100439 4, HEMBA1004396, HEMBA1004429, HEMBA100460, HEMBA1004617, HEMBA1004611, HEMBA1004631, HEMBA1004631, HEMBA1004670, HEMBA1004733, HEMBA1004816, HEMBA1004820, HEMBA1004918, HEMBA1004960, HEMBA1004978, HEMBA1004995, HEMBA1005008, HEMBA1005062, HEMBA1005293, HEMBA1005304, HEMBA1005315, HEMBA1005372, HEMBA1005389, HEMBA1005570, HEMBA1005616, HEMBA1005627, HEMBA1005631, HEMBA1005632, HEMBA1005670, HEMBA1005699, HEMBA1005705, HEMBA1005765, HEMBA1005780, HEMBA1005853, HEMBA1005909, HEMBA1005909, HEMBA1005705, HEMBA1005765, HEMBA1005780, HEMBA1005853, HEMBA1005909,

HEMBA1005931, HEMBA1005934, HEMBA1005962, HEMBA1005999, HEMBA1006002, HEMBA1006005, HEMBA1006042, HEMBA1006067, HEMBA1006090, HEMBA1006142, HEMBA1006268, HEMBA1006334, HEMBA1006359, HEMBA1006416, HEMBA1006419, HEMBA100642 1, HEMBA1006446, HEMBA1006471, HEMBA1006486, HEMBA1006489, HEMBA1006546, HEMBA10065 69.

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HEMBA1006849, HEMBA1006865, HEMBA1006929, HEMBA1007045, HEMBA1007051, HEMBA100 7073,
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HEMBB1000643, HEMBB1000665, HEMBB1000726, HEMBB1000738, HEMBB10 00770, HEMBB1000794,
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NT2RP3001527, NT2RP3001607, NT2RP3001752, NT2RP3001782, NT2RP3001926, NT2RP3001 989,
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NT2RP3003746, NT2RP3003799, NT2RP3003989, NT2RP3004016, NT2RP3004095, NT2RP3 004110,
NT2RP3004145, NT2RP3004148, NT2RP3004215, NT2RP3004399, NT2RP3004475, NT2RP 3004503,
NT2RP4000321, NT2RP4000519, NT2RP4000996,
OVARC1000133, OVARC1000198, OVARC1000240, OVARC1000427, OVARC1000431, OVARC1000533 ,
OVARC1000543, OVARC1000573, OVARC1000578, OVARC1000678, OVARC1000679, OVARC100070 0.
OVARC1000769, OVARC1000802, OVARC1000891, OVARC1000897, OVARC1000936, OVARC10009 37,
OVARC1000960, OVARC1000971, OVARC1001000, OVARC1001062, OVARC1001072, OVARC1001 085,
OVARC1001118, OVARC1001129, OVARC1001169, OVARC1001261, OVARC1001268, OVARC100 1282,
OVARC1001330, OVARC1001339, OVARC1001342, OVARC1001357, OVARC1001442, OVARC10 01480,
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OVARC1001901, OVARC1001911, OVARC1001989, OVARC1002044, OVARC1002066, OVARC 1002143,
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PLACE1000421, PLACE1000424, PLACE1000481, PLACE1000749, PLACE1000841, PLACE100093 1,
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PLACE1006521, PLACE1006534, PLACE1006540, PLACE1006617, PLACE1006640, PLACE1 006779,
PLACE1006792, PLACE1006800, PLACE1006805, PLACE1006815, PLACE1006860, PLACE 1006878,
PLACE1007097, PLACE1007132, PLACE1007140, PLACE1007276, PLACE1007286, PLAC E1007525,
PLACE1007677, PLACE1007737, PLACE1007866,
PLACE1008045, PLACE1008111, PLACE1008201, PLACE1008231, PLACE1008392, PLACE1008424,
PLACE1008437, PLACE1008455, PLACE1008625, PLACE1008748, PLACE1008757, PLACE100879 8,
PLACE1008807, PLACE1008920, PLACE1008947, PLACE1009150, PLACE1009246, PLACE10093 08,
PLACE1009410, PLACE1009477, PLACE1009613, PLACE1009639, PLACE1009879, PLACE1009 888,
PLACE1009924, PLACE1009935, PLACE1010069, PLACE1010083, PLACE1010181, PLACE101 0341,
PLACE1010624, PLACE1010628, PLACE1010631, PLACE1010662, PLACE1010916,
                                                                                                       PLACE10 10947,
PLACE1011057, PLACE1011133, PLACE1011143, PLACE1011296, PLACE1011340,
                                                                                                       PLACE1 011419.
PLACE1011503, PLACE1011641, PLACE1011649, PLACE1011650, PLACE1011729, PLACE 1011778,
PLACE1011874, PLACE1011962, PLACE1011995,
PLACE2000003, PLACE2000103, PLACE2000115, PLACE2000132, PLACE2000140, PLACE2000176,
PLACE2000317, PLACE2000342, PLACE2000419, PLACE2000435, PLACE2000455, PLACE200046 5,
PLACE3000004, PLACE3000124, PLACE3000136, PLACE3000158, PLACE3000169, PLACE30001 99,
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PLACE3000208, PLACE3000218, PLACE3000341, PLACE3000362, PLACE3000365, PLACE3000 388, PLACE3000400, PLACE3000402, PLACE3000475, PLACE4000093, PLACE4000100, PLACE400 0233, PLACE4000252, PLACE4000379, PLACE4000411, PLACE4000494, THYRO1000026, THYRO1000035, THYRO1000085, THYRO1000132, THYRO1000156, THYRO1000186, THYRO1000190, THYRO1000221, THYRO1000241, THYRO1000438, THYRO1000452, THYRO100055 8, THYRO1000602, THYRO1000641, THYRO1000658, THYRO1000699, THYRO1000748, THYRO10007 96,

THYRO1000843, THYRO1000895, THRO1000974, THYRO1000975, THYRO1001062, THYRO1001 093, THYRO1001121, THYRO1001177, THYRO1001262, THYRO1001290, THYRO1001322, THYRO100 1411, THYRO1001480, THYRO1001541, THYRO1001573, THYRO1001617, THYRO1001895, Y79AA10 00065, Y79AA1000405, Y79AA1000538, Y79AA1000802, Y79AA1000969, Y79AA1001061, Y79AA1 001167, Y79AA1001384, Y79AA1001594, Y79AA1001875, Y79AA1002211,

Group (11): 228 clones

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The following 228 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according
to the fullness ratio shown in Table 7. The clones were a novel clone, in which at least the 3'-end sequence was not
identical to any of human ESTs.
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HEMBA1001979, HEMBA1002160, HEMBA1002162, HEMBA1002166, HEMBA1002389, HEMBA1002 498,
HEMBA1002592, HEMBA1002679, HEMBA1002944, HEMBA1003034, HEMBA1003142, HEMBA100 3548,
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HEMBB1001967, HEMBB1002043, HEMBB1002094, HEMBB1002190, HEMBB1002520, HEMBB100 2556,
HEMBB1002590.
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MAMMA1000431, MAMMA1000483, MAMMA1000559, MAMMA1000723, MAMMA1000744, MAMMA100075 2,
MAMMA1000782, MAMMA1000802, MAMMA1000851, MAMMA1000877, MAMMA1000962, MAMMA10010 74,
MAMMA1001133, MAMMA1001191, MAMMA1001206, MAMMA1001249, MAMMA1001330, MAMMA1001 452,
MAMMA1001502, MAMMA1001663, MAMMA1001806, MAMMA1001836, MAMMA1001880, MAMMA100 1890,
MAMMA1001908, MAMMA1001963, MAMMA1002058, MAMMA1002155, MAMMA1002158, MAMMA10 02230,
MAMMA1002282, MAMMA1002293, MAMMA1002332, MAMMA1002359, MAMMA1002494, MAMMA1 002573,
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MAMMA1003140.
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PLACE1005085, PLACE1005666, PLACE1005845, PLACE1006164, PLACE1006262, PLACE10067 54,
PLACE1007852, PLACE1008331, PLACE1008643, PLACE1009050, PLACE1009155, PLACE1009 172,
PLACE1009183, PLACE1009200, PLACE1009398, PLACE1009595, PLACE1010702, PLACE101 0891,
PLACE1011165, PLACE1011203, PLACE1011375, PLACE1011576, PLACE1011762, PLACE10 11964, PLACE2000136, PLACE2000264, PLACE2000305, PLACE2000379, PLACE3000220, PLACE4 000250,
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THYRO1000092,THYRO1000471, THYRO1000852, THYRO1001534, THYRO1001595, THYRO 1001745.

## **EXAMPLE 12**

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Homology search using the 5'-end and 3'-end sequences of the selected clones.

<sup>5</sup> [0172] The 5'-end sequences of the selected 5547 clones were used for the homology search of the SwissProt, and both the 5'-end 3'-end sequences were used for the search of the GenBank and UniGene (ref. the result of the search of the SwissProt, GenBank (except ESTs and STSs), and UniGene (Human) was attached).

[0173] Each search result is shown in the last part of this SPECIFICATION by arranging each item in the following format.

		5'-end sequence	3'-end sequence
	Swiss-Prot	Homology search result 1	****
	GenBank	Homology search result 2	Homology search result 3
15	UniGene	Homology search result 4	Homology search result 5

[0174] According to the top hit data, at least 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth. Among the clones predicted belonging to any of the categories, 1001 clones were estimated to have a relatively high homology with the known proteins or genes in the same category. In addition, 429 clones were estimated to have a relatively low homology with the known proteins in the same category. [0175] Herein, the term "relatively high homology" is defined as having 60% or more identity and the P-value 10<sup>-10</sup> or less in comparison with known sequences in the SwissProt database, or 64% or more identity and the P-value 10<sup>-15</sup> or less in comparison with those in the GenBank and UniGene databases (see the attached list). Also, the term "relatively low homology" is defined as not fulfilling the requirements to be "relatively high homology", but still having the scores, 25% or more identity and the P-value 10-6 or less, using the sequence having 55 nucleotides or more, in comparison with known sequences in the SwissProt database (see the attached list). The P-value is a score obtained statistically by taking into account the probability of occurrence of the similarity between two sequences. In general, the smaller P-value reflects the higher similarity (Altschul S.F., Gish W., Miller W., Myers E.W., and Lipman D.J. (1990) "Basic local alignment search tool" J.Mol. Biol., 215: 403-410; Gish W., and States D.J. (1993) "Identification of protein coding regions by database similarity search" Nature Genet. 3: 266-272).

[0176] The clones predicted to encode a protein in the category of secretory protein or membrane protein have the keywords, "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", or "connective tissue", or descriptions from which the clone can be predicted to be a secretory or membrane protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0177] The clones predicted to encode a protein in the category of glycoprotein have the keywords, "glycoprotein", or descriptions from which the clone can be predicted to be a glycoprotein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence. [0178] The clones predicted to encode a protein in the category of proteins associated with signal transduction have the keywords, "serine/threonine-protein kinase", "tyrosine-protein kinase", "SH3 domain", or "WD repeat", or descriptions from which the clone can be predicted to be a protein associated with signal transduction (such as "ADP-ribosylation factor"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0179] The clones predicted to encode a protein in the category of proteins associated with transcription have the keywords, "transcription regulation", "zinc finger", or "homeobox", or descriptions from which the clone can be predicted to be a protein associated with transcription, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0180] The clones predicted to encode a protein in the category of proteins associated with diseases are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein that is deposited in the Online Mendelian Inheritance in Man (OMIM) database, which is a database of human genes and diseases, or the top hit data has descriptions from which the clone can be predicted to be a protein associated with diseases.

[0181] The clones predicted to encode a protein in the category of enzyme or proteins associated with metabolism are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein with E.C.No. (Enzyme commission number), or the top hit data has descriptions from which the clone can be predicted to be an enzyme or protein associated with metabolism (such as "metabolism", "oxidoreductase", or "lipid").

[0182] The clones predicted to encode a protein in the category of proteins associated with cell division or cell proliferation have the keywords, "cell division", "cell cycle", "mitosis", or "chromosomal protein", or descriptions from which the clone can be predicted to be a protein associated with cell division or cell proliferation (such as "histone", "cell growth", or "apoptosis"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0183] The clones predicted to encode a protein in the category of proteins associated with cytoskeleton have the keywords, "structural protein", "cytoskeleton", "actin-binding", or "microtubules", or descriptions from which the clone can be predicted to be a protein associated with cytoskeleton, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0184] The clones predicted to encode a protein in the category of proteins associated with RNA synthesis include the above clones predicted to be a protein associated with transcription, and also the clones which have the keywords, "RNA splicing", or "RNA processing", or descriptions from which the clone can be predicted to be a protein associated with RNA synthesis (such as "RNA helicase", "polyadenylation", or "RNA transport"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0185] The clones predicted to encode a protein in the category of nuclear protein include the above clones predicted to be a protein associated with transcription, and also the clones which have the keyword, "nuclear protein", or descriptions from which the clone can be predicted to be a nuclear protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0186] The clones predicted to encode a protein in the category of proteins associated with protein synthesis or transport have the keywords, "translation regulation", "protein biosynthesis", "amino-acid biosynthesis", "ribosomal protein", or "protein transport", or descriptions from which the clone can be predicted to be a protein associated with protein synthesis or transport (such as "signal recognition particle", "ubiquitin", "proteosome", or "protease"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0187] The clones predicted to encode a protein in the category of proteins associated with cellular defense have the keywords, "heat shock", "chaperone", "DNA repair", or "DNA damage", or descriptions from which the clone can be predicted to be a protein associated with cellular defense, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0188] The clones predicted to encode a protein in the category of proteins associated with development or growth have the keyword, "developmental protein", or descriptions from which the clone can be predicted to be a protein associated with development or growth, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0189] The following 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth.

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HEMBA100005, HEMBA100012, HEMBA100020, HEMBA100030, HEMBA1000158, HEMBA1000185, HEMBA1000211, HEMBA1000216, HEMBA1000303, HEMBA1000459, HEMBA1000488, HEMBA100049 1, HEMBA1000523, HEMBA1000531, HEMBA1000542, HEMBA1000561, HEMBA1000569, HEMBA10005 88, HEMBA1000852, HEMBA1000972, HEMBA1000991, HEMBA1000752, HEMBA1000972, HEMBA1000991, HEMBA1001017, HEMBA1001017, HEMBA1001088, HEMBA1001123, HEMBA1001137, HEMBA1001174, HEMBA100 01257, HEMBA1001302, HEMBA1001351, HEMBA1001387, HEMBA1001407, HEMBA1001476, HEMBA1 001510, HEMBA1001579, HEMBA1001579, HEMBA1001595, HEMBA1001620, HEMBA 1001672, HEMBA1001819, HEMBA1001714, HEMBA1001744, HEMBA1001809, HEMBA1001819, HEMBA1001822, HEMBA1001847, HEMBA1001896, HEMBA1002035, HEMBA1002035, HEMBA1002021, HEMBA1002229, HEMBA1002557, HEMBA1002361, HEMBA1002417, HEMBA1002495, HEMBA1002513, HEMBA1002510, HEMBA1002555, HEMBA1002569, HEMBA1002609, HEMBA1002688, HEMBA1002716, HEMBA1002810, HEMBA1002896,
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                                Y79AA1000705,
                                                Y79AA1000734,
                                                                Y79AA1000752,
                                                                                Y79AA1000784.
Y79AA1000833,
                Y79AA1000966,
                                Y79AA1000968,
                                                Y79AA1000985.
                                                                Y79AA1001105,
                                                                                Y79AA1001236,
Y79AA1001299,
                Y79AA1001391.
                                Y79AA1001533,
                                                Y79AA1001548,
                                                                Y79AA1001679,
                                                                                Y79AA1001827,
Y79AA1001866, Y79AA1002093, Y79AA1002258, Y79AA1002361, Y79AA1002416,
[0191] The following 429 clones were estimated to have a relatively low homology with the known proteins in the
same category.
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	HEMBA1000012, HEMBA1000752,	HEMBA1000185,	HEMBA1000488,	HEMBA1000491,	HEMBA1000569,
5		НЕМВА1001407, НЕМВА1001809, НЕ		HEMBA1001744,	HEMBA1001800,
10		НЕМВА1002102, НЕМВА1002609, НЕ		HEMBA1002212, A1002939,	HEMBA1002495,
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	· · · · · · · · · · · · · · · · · · ·	HEMBA1003077, HEMBA1003136, HEMBA1003179, HEMBA1003235, HEMBA1003418,
5	HEMBA1003568, HEMBA1003953,	HEMBA1003598, HEMBA1003680, HEMBA1003684, HEMBA1003836,
10	HEMBA1004276,	HEMBA1004408, HEMBA1004669, HEMBA1004734,
	HEMBA1005475,	НЕМВА1005595, НЕМВА1005737, НЕМВА1005815, НЕМВА1005931,
15		HEMBA1006474, HEMBA1006521, HEMBA1006562, HEMBA1006652, HEMBA1006914,
	HEMBA1007045,	НЕМВА1007121, ЏЕМВА1007300, НЕМВА1007301,
20	HEMBB1000119,	HEMBB1000217, HEMBB1000317, HEMBB1000632, HEMBB1000789,
	HEMBB1001011,	HEMBB1001177, HEMBB1001234, HEMBB1001282, HEMBB1001288,
25	HEMBB1002050,	HEMBB1002092,
		MAMMA1000672, MAMMA1000841, MAMMA1000897, MAMMA1001008, MAMMA1001041,
30	·	MAMMA1002573, MAMMA1002619, MAMMA1002636, MAMMA1002650, MAMMA1002881, MAMMA1002937,
35	NT2RM1000118,	NT2RM1000153, NT2RM1000186, NT2RM1000187, NT2RM1000377,
	· ·	NT2RM1000666, NT2RM1000691, NT2RM1000770, NT2RM1000772, NT2RM1000882, NT2RM1000898,
40	NT2RM1001059,	NT2RM1001072, NT2RM1001139,
	NT2RM2000363, NT2RM2000623,	NT2RM2000371, NT2RM2000402, NT2RM2000577, NT2RM2000599, NT2RM2000714,
45	NT2RM2001131,	NT2RM2001221, NT2RM2001324, NT2RM2001345,
50	NT2RM2001588, NT2RM2001785, NT2RM2002580,	NT2RM2001635, NT2RM2001670, NT2RM2001700, NT2RM2001730, NT2RM2001797, NT2RM2001823, NT2RM2001936, NT2RM2001989,
55	NT2RM4000030, NT2RM4000191, NT2RM4000734,	NT2RM4000085, NT2RM4000155, NT2RM4000156, NT2RM4000169, NT2RM4000215, NT2RM4000354, NT2RM4000421, NT2RM4000496, NT2RM4000751,

5	NT2RM4001032, NT2RM4001444,	NT2RM4001140,	NT2RM4001316,	NT2RM4001340,	NT2RM4001410,
•	NT2RM4001611, NT2RM4001930,	NT2RM4001629, NT2RM4001979, NT2	NT2RM4001746, 2RM4001987,	NT2RM4001776,	NT2RM4001783,
10	NT2RM4002062,	NT2RM4002145, NT2	2RM4002189, NT2R	M4002226, NT2RM40	002409,
	NT2RM4002527,	NT2RM4002571, NT2	ZRM4002623,		
15	NT2RP1000111, NT2RP1000460,	NT2RP1000130, NT2RP1000470, NT2	NT2RP1000333, 2RP1000493,	NT2RP1000348,	NT2RP1000363,
20	NT2RP1000547, NT2RP1001543,	NT2RP1000609,	NT2RP1001033,	NT2RP1001080,	NT2RP1001313,
	NT2RP2000006, NT2RP2000448,	NT2RP2000133,	NT2RP2000161,	NT2RP2000183,	NT2RP2000257,
25	NT2RP2000710, NT2RP2001174,	NT2RP2000764, NT2RP2001839, NT2	NT2RP2000812, CRP2001900,	NT2RP2000932,	NT2RP2001127,
30	NT2RP2002025, NT2RP2002618,	NT2RP2002058, NT2RP2002710, NT2	NT2RP2002185, 2RP2002980,	NT2RP2002442,	NT2RP2002595,
	NT2RP2003125, NT2RP2003564,	NT2RP2003277, NT2RP2003706,	NT2RP2003286,	NT2RP2003293,	NT2RP2003506,
35	NT2RP2004098,	NT2RP2004187.	NT2RP2004568,	NT2RP2004791,	NT2RP2005022,
	NT2RP2005037,	NT2RP2005038,	NT2RP2005204,	NT2RP2005344,	NT2RP2005465,
	NT2RP2005496,	NT2RP2005531,	NT2RP2005690,	NT2RP2005763,	NT2RP2005767,
40	NT2RP2006275,				•
	NT2RP3000046,	NT2RP3000047,	NT2RP3000361,	NT2RP3000397,	NT2RP3000418,
	NT2RP3000527,	NT2RP3000590,	NT2RP3000603,	NT2RP3000759,	NT2RP3000868,
45	NT2RP3000994,				
	NT2RP3001216,	NT2RP3001274,	NT2RP3001355.	NT2RP3001399,	NT2RP3001407,
	<u>-</u>	NT2RP3001427,			,
50	NT2RP3001587, NT2RP3002399,	NT2RP3001708, NT2RP3002663, NT2	NT2RP3002056, RP3002785, NT2R		NT2RP3002377,
55	NT2RP3003061, NT2RP3003825,	NT2RP3003500, NT2RP3003831,	NT2RP3003672,	NT2RP3003701,	NT2RP3003716,

			•		
	NT2RP3004209,	NT2RP3004253,	NT2RP3004454,	NT2RP3004472,	NT2RP3004475,
	NT2RP3004507	NT2RP3004569, NT			
5			•	•	,
	NT2RP4000210,	NT2RP4000212,	NT2RP4000259,	NT2RP4000312,	NT2RP4000370,
	NT2RP4000417,	•	NT2RP4000457,	NT2RP4000498,	NT2RP4000528,
	NT2RP4000879	NT2RP4000925,		,	
10		,			
	NT2RP4001041,	NT2RP4001095,	NT2RP4001122,	NT2RP4001126,	NT2RP4001148,
	NT2RP4001149,	•	NT2RP4001174,	NT2RP4001207,	NT2RP4001219,
	NT2RP4001372,	NT2RP4001375,	,	,	
15					
	NT2RP4001551,	NT2RP4001592,	NT2RP4001638,	NT2RP4001725,	NT2ŔP4001790,
	NT2RP4001823,	•	NT2RP4001849,	NT2RP4001893,	NT2RP4001927,
	•	NT2RP4002408,	,	,	
20		•			
20	NT2RP5003477,	NT2RP5003522, NT	2RP5003524.		_
		•	•		•
	OVARC1000004,	OVARC1000087,	OVARC1000139,	OVARC1000309,	OVARC1000543,
	OVARC1000746,	OVARC1000846, OV	•	•	
25				•	
	OVARC1001180,	OVARC1001329,	OVARC1001476,	OVARC1001555,	OVARC1001610,
	OVARC1001762,	OVARC1001880, OV	ARC1001942, OVAR		
30	PLACE1000066,	PLACE1000142,	PLACE1000420,	PLACE1000547,	PLACE1000583,
	PLACE1000863,				
	PLACE1001387,	PLACE1001632, PL	ACE1001692, PLAC	E1001989,	
35					
	PLACE1002450,	PLACE1002571,	PLACE1002722,	PLACE1002775,	PLACE1002811,
	PLACE1002991,				
		•			•
40	PLACE1003190,	PLACE1003258,	PLACE1003420,	PLACE1003493,	PLACE1003605,
	PLACE1003669,	PLACE1003704, PL	ACE1003915,		
	PLACE1004197,	PLACE1004428,	PLACE1004471,	PLACE1004629,	PLACE1004751,
45	PLACE1004902,				
	DI 467144554	<b>DI</b> ( <b>C C C C C C C C C C</b>			
	PLACE1005287,	PLACE1005557, PLA	ACE1005611, PLAC	E1005763, PLACE1	005890,
	DI 100100000	DI 400400000			
50	PLACE1006157,	PLACE1006469, PLA	CE1006731,		
	DI LODI CORCO	D1 4-D4			
	PLACE1007238,		PLACE1007488,	PLACE1007544,	PLACE1007598,
	PLACE1007729,	PLACE1008244, PLA	CE1008275, PLAC	E1008947,	
55	PLACE1009094.	PLACE1009319,			
		DI ACCIONOSO	PLACE1009622,	PLACE1009721,	PLACE1009845,

PLACE1009861,

5	PLACE1010083,	PLACE1010089,	PLACE1010106,	PLACE1010152,	PLACE1010194,
	PLACE1010310,	PLACE1010833,	PLACE1010960,	PLACE1011056,	PLACE1011219,
	PLACE1011263,	PLACE1011371, PL	ACE1011492,		

- PLACE2000034, PLACE2000164, PLACE2000274, PLACE2000399, PLACE2000411, PLACE2000458,
- PLACE3000148, PLACE3000156, PLACE3000242, PLACE3000322, PLACE3000353, PLACE3000373,
  - PLACE4000009, PLACE4000014, PLACE4000259, PLACE4000431, PLACE4000522, PLACE4000558, PLACE4000581, PLACE4000590, PLACE4000612,

SKNMC1000011,

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THYR01000072, THYR01000852, THYR01000934, THYR01000951,

Y79AA1000258, Y79AA1000368, Y79AA1000539, Y79AA1000589, Y79AA1000962; Y79AA1001048, Y79AA1001233, Y79AA1001394, Y79AA1001581, Y79AA1001613, Y79AA1001692, Y79AA1001848, Y79AA1001874, Y79AA1001963,

Y79AA1002027, Y79AA1002139, Y79AA1002208, Y79AA1002209, Y79AA1002472, Y79AA1002482,

(1) Among the 1430 clones, the following 259 clones were predicted to encode a protein in the category of secretory protein or membrane protein (including the clones belonging to plural categories).

HEMBA1000005, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001351, HEMBA1001407, HEMBA1001476, HEMBA1001569, HEMBA1001678, HEMBA1001804, HEMBA1002150, HEMBA100254 7, 40 HEMBA1002555, HEMBA1002688, HEMBA1002716, HEMBA1002999, HEMBA1003071, HEMBA10030 77, HEMBA1003538, HEMBA1003598, HEMBA1003690, HEMBA1004207, HEMBA1004356, HEMBA1004 389, HEMBA1004756, HEMBA1005338, HEMBA1005367, HEMBA1005576, HEMBA1005699, HEMBA100 5963, HEMBA1005990, HEMBA1006310, HEMBA1006344, HEMBA1006562, HEMBA1006976, HEMBA10 07045, HEMBA1007301, HEMBB1000037, HEMBB1000119, HEMBB1000317, HEMBB1000530, HEMBB1 000915, 45 HEMBB1001112, HEMBB1001119, HEMBB1001151, HEMBB1001282, HEMBB1001331, HEMBB 1002042, HEMBB1002092, HEMBB1002193, HEMBB1002600, MAMMA1000103, MAMMA1000424, MAMM A1000672, MAMMA1000734, MAMMA1000897, MAMMA1001008, MAMMA1001030, MAMMA1001041, MAM MA1001075, MAMMA1001080, MAMMA1001751, MAMMA1001754, MAMMA1002143, MAMMA1002219, MA MMA1002485, MAMMA1002636, MAMMA1002938, MAMMA1003104, NT2RM1000199, NT2RM1000355, N T2RM1000430, 50 NT2RM1000648, NT2RM1000742, NT2RM1000770, NT2RM1000811, NT2RM1000833, NT2RM1000867, NT2RM1000882, NT2RM1000883, NT2RM1001139, NT2RM2000374, NT2RM2000402, NT2RM2000422, NT2RM2000566, NT2RM2000623, NT2RR2001324, NT2RM2001499, NT2RM200158 8, NT2RM2001592, NT2RM2001613, NT2RM2001635, NT2RM2001648, NT2RM2001760, NT2RM20021 09, NT2RM2002145, NT2RM4000156, NT2RM4000169, NT2RM4000386, NT2RM4000433, NT2RM4000795, NT2RM4001032, 55 NT2RM4001054, NT2RM4001410, NT2RM4001605; NT2RM4001930, NT2RM4001987, NT2RM4002073, NT2RM4002145, NT2RM4002189, NT2RM4002558, NT2RM400256 5, NT2RM4002571, NT2RM4002594, NT2RP1000130, NT2RP1000326, NT2RP1000363, NT2RP10004 13, NT2RP1000547, NT2RP1000782, NT2RP1000856, NT2RP1001247, NT2RP1001313, NT2RP1001 546, NT2RP1001569, NT2RP2000056,

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NT2RP2000257, NT2RP2000842, NT2RP2001081, NT2RP200 1991, NT2RP2002025, NT2RP2002193,
       NT2RP2002256, NT2RP2002312, NT2RP2002479, NT2RP20 02504, NT2RP2003433, NT2RP2003706,
       NT2RP2003760, NT2RP2004194, NT2RP2004655, NT2RP2 005020, NT2RP2005344, NT2RP2005465,
       NT2RP2005752, NT2RP2005753, NT2RP2005835, NT2RP 2005933, NT2RP2006275, NT2RP2006565,
5
       NT2RP3006571, NT2RP3000324, NT2RP3000739, NT2R P3000919, NT2RP3001140, NT2RP3001355,
       NT2RP3001676, NT2RP3001708, NT2RP3001727, NT2 RP3002659, NT2RP3003059, NT2RP3003101,
       NT2RP3003203, NT2RP3003212, NT2RP3003242, NT 2RP3003672, NT2RP3003701, NT2RP3003716,
       NT2RP3003918, NT2RP3004207, NT2RP3004253, N T2RP4000008, NT2RP4000212, NT2RP4000243,
       NT2RP4000417, NT2RP4000878, NT2RP4000907, NT2RP4000925, NT2RP4000928, NT2RP4001079,
10
       NT2RP4001117, NT2RP4001150, NT2RP4001174, NT2RP4001313, NT2RP4001315, NT2RP4001345,
       NT2RP4001372, NT2RP4001574, NT2RP400182 3, NT2RP4001966, NT2RP5003524, OVARC1000060,
       OVARC1000326, OVARC1000543, OVARC10006 82, OVARC1000722, OVARC1000912, OVARC1001055,
       OVARC1001154, OVARC1001261.
       OVARC1001329, OVARC1001419, OVARC1001506, OVARC1001542, OVARC1001610, OVARC1001713,
15
       OVARC1001802, OVARC1001943, PLACE1000611, PLACE1001092, PLACE1001136, PLACE100129 4,
       PLACE1002437, PLACE1002500, PLACE1002722, PLACE1002782, PLACE1002816, PLACE10034 20,
       PLACE1003493, PLACE1003596, PLACE1004104, PLACE1004197, PLACE1004203, PLACE1004 277,
       PLACE1004629, PLACE1004751, PLACE1005494, PLACE1005804, PLACE1005890, PLACE100 6157,
       PLACE1007409, PLACE1007852, PLACE1008044, PLACE1008273, PLACE1008309, PLACE10 08603,
20
       PLACE1008643, PLACE1009659, PLACE1009670, PLACE1009845, PLACE1009861, PLACE1 011371,
       PLACE1011492, PLACE1011896, PLACE2000034, PLACE2000341, PLACE2000399, PLACE 2000458,
       PLACE3000020, PLACE3000322, PLACE3000353, PLACE4000052, PLACE4000269, PLAC E4000522,
       PLACE4000581, THYRO1000783, Y79AA1000258, Y79AA1000346, Y79AA1001874,
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<sup>25</sup> (2) The following 111 clones were predicted to encode a protein in the category of glycoprotein (including the clones belonging to plural categories).

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HEMBA1000752, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001302, HEMBA1002150, HEMBA1002547, HEMBA1002555, HEMBA1003077, HEMBA1003538, HEMBA1003598, HEMBA100386 6, HEMBA1004356, HEMBA1005338, HEMBA1005576, HEMBA1005699, HEMBA1006562, HEMBA10069 76, HEMBA1007301, HEMBB1000317, HEMBB1000530, HEMBB1001119, HEMBB1002092, HEMBB1002 193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388, MAMMA1001771, MAMMA100 2485, MAMMA1002636, MAMMA1002938, NT2RM1000648, NT2RM2000374, NT2RM2000407, NT2RM20 00422, NT2RM2000623, NT2RM2001499, NT2RM2001588, NT2RM2001839, NT2RM2001930, NT2RM4 000156, NT2RM4000233, NT2RM4001410, NT2RM4001987, NT2RM4002145, NT2RM4002189, NT2RM 4002194, NT2RM4002558, NT2RM4002571, NT2RP1000856, NT2RP1001247, NT2RP2000056, NT2R P2001991, NT2RP2002025, NT2RP2002325, NT2RP2002385, NT2RP2003706, NT2RP2005933, NT2 RP3001140, NT2RP3003242, NT2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003914, NT 2RP3004253, NT2RP4000212, NT2RP4000417, NT2RP4000925, NT2RP4001149, NT2RP4001150, N T2RP4001219, NT2RP4001345, NT2RP4001372, NT2RP5003522, NT2RP5003524, OVARC1000543, OVARC1000682, OVARC1000722, OVARC1001055, OVARC1001154, OVARC1001506, OVARC1001713 PLACE1002437, PLACE1002722, PLACE1003258, PLACE1003493, PLACE1004197, PLACE100475 1, PLACE1005698, PLACE1005804, PLACE1006157, PLACE1006754, PLACE1008643, PLACE10090 94, PLACE1009861, PLACE1011371, PLACE2000034, PLACE2000062, PLACE2000399, PLACE2000458, PLACE3000009, PLACE3000020, PLACE3000353, PLACE3000373, PLACE4000052, PLACE4000230, PLACE4000522, PLACE4000581, THYRO1000852, Y79AA1001874,

(3) The following 147 clones were predicted to encode a protein in the category of proteins associated with signal transduction (including the clones belonging to plural categories).

HEMBA100030, HEMBA1000185, HEMBA1000303, HEMBA1000459, HEMBA1000491, HEMBA1000657, HEMBA1000991, HEMBA1001174, HEMBA1001387, HEMBA1001822, HEMBA1001921, HEMBA100221 2, HEMBA1002341, HEMBA1002417, HEMBA1002810, HEMBA1002896, HEMBA1003291, HEMBA10033 14, HEMBA1003560, HEMBA1003615, HEMBA1003621, HEMBA1003805, HEMBA1004286, HEMBA1004 604, HEMBA1006173, HEMBA1006540, HEMBA1006973, HEMBA1007174, HEMBB1000632, HEMBB100 0781, HEMBB1001051, HEMBB1001177, HEMBB1001234, HEMBB1001294, HEMBB1001384, HEMBB10 02193, HEMBB1002477, HEMBB1002635, MAMMA1001305, MAMMA1002699, MAMMA1002842, MAMMA1 003057, MAMMA1003113, NT2RM1000118, NT2RM1000186, NT2RM1000772, NT2RM1000850, NT2RM 1001072, NT2RM2000124, NT2RM2000612, NT2RM2001065, NT2RM2001221, NT2RM2001345, NT2R M2001652, NT2RM2001983, NT2RM2002109, NT2RM2002128, NT2RM4000327, NT2RM4000354, NT2 RM4000798,

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NT2RM4001203, NT2RM4001411, NT2RM4001412, NT2RM4001582, NT2RM4001629, NT 2RM4001758, NT2RM4002226, NT2RP1000018, NT2RP1000112, NT2RP1000701, NT2RP1000825, N T2RP1001395, NT2RP1001410, NT2RP1001457, NT2RP1001482, NT2RP2002058, NT2RP2002710, NT2RP2003702, NT2RP2003912, NT2RP2004232, NT2RP2004768, NT2RP2005022, NT2RP2005620, NT2RP2005890, NT2RP3000299, NT2RP3000403, NT2RP3000742, NT2RP3000845, NT2RP300176 4, NT2RP3002004, NT2RP3002785, NT2RP3002999, NT2RP3003230, NT2RP3003411, NT2RP30034 64, NT2RP3003800, NT2RP3004475, NT2RP3004534, NT2RP4000147, NT2RP4000376, NT2RP4001122, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001760, NT2RP4001849, NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000437, OVARC1000465, OVARC100064 9, OVARC1000945, OVARC1001476, OVARC1001861, PLACE1001377, PLACE1001384, PLACE10013 87, PLACE1002591, PLACE1002685, PLACE1003353, PLACE1004128, PLACE1004 918, PLACE1007488, PLACE1005739, PLACE1005968, PLACE1006385, PLACE1007238, PLACE100 7375, PLACE1007488, PLACE1008244, PLACE1008650, PLACE1009319, PLACE1009468, PLACE10 10083, PLACE1010942, PLACE1011041, PLACE1011046, PLACE1011923, PLACE2000164, PLACE3 000145, PLACE3000350, PLACE3000475, THYRO1001595, Y79AA1000059, Y79AA1000966,
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(4) The following 230 clones were predicted to encode a protein in the category of proteins associated with transcription (including the clones belonging to plural categories).

HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000851, HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA100184 7, HEMBA1002035. HEMBA1002092. HEMBA1002569, HEMBA1002609, HEMBA1003545, HEMBA10035 68, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004 353, HEMBA1004479, HEMBA1004758, HEMBA1005359, HEMBA1005528, HEMBA1005548, HEMBA100 5931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006559, HEMBB1000250, HEMBB10 00789, HEMBB1001011, HEMBB1001314, HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1 002217, MAMMA1000155, MAMMA1000183, MAMMA1000388, MAMMA1001633, MAMMA1001820, MAMMA 1001837, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000666, NT2RM1000691, NT2R M1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2 RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2002580, NT2RM4000024, NT 2RM4000104, NT2RM4000191, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, N T2RM4000996, NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RP1000111, NT2RP1000333, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP2000008, NT2RP200013 3, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2002252, NT2RP20025 03, NT2RP2002591,

NT2RP2003125, NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004187, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP2006238, NT2RP200646 4, NT2RP3000050, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP30006 32, NT2RP3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001 399, NT2RP3001671, NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP300 1855, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP30 03251, NT2RP3003327, NT2RP3004258, NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3 004617, NT2RP4000078, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000787, NT2RP 4000865, NT2RP4000997, NT2RP4001029, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2R P4001551, NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVA RC1000479, OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577, OVARC1001949, PL ACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706, PLACE1001118, P LACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, PLACE1003302, PLACE1003605, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004681, PLACE1006414, PLACE1006438, PLACE1006482, PLACE100723 9, PLACE1007346, PLACE1007544, PLACE1007598, PLACE1007955, PLACE1007969, PLACE10086 27, PLACE1008941,

PLACE1009099, PLACE1009246, PLACE1009398, PLACE1010261,
PLACE1010491, PLACE1010702, PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072,
PLACE2000359, PLACE4000128, PLACE4000156, PLACE4000259, PLACE4000431, THYRO100012 1,
THYRO1000242, THYRO1000501, THYRO1000585, THYRO1001100, THYRO1001189, Y79AA10000 37,
Y79AA1000627, Y79AA1000705, Y79AA1001105, Y79AA1001299, Y79AA1001391, Y79AA1001 533,
Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1002093, Y79AA1002472, Y79AA100 2482,

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(5) The following 436 clones were predicted to encode a protein in the category of proteins associated with diseases
        (including the clones belonging to plural categories).
        HEMBA1000158, HEMBA1000216, HEMBA1000523, HEMBA1000561, HEMBA1000569, HEMBA1000673,
        HEMBA1000972, HEMBA1001017, HEMBA1001019, HEMBA1001059, HEMBA1001071, HEMBA100108 8,
        HEMBA1001123, HEMBA1001407, HEMBA1001569, HEMBA1001570, HEMBA1001672, HEMBA10020 35,
        HEMBA1002160, HEMBA1002162, HEMBA1002547, HEMBA1002939, HEMBA1003077, HEMBA1003 418,
        HEMBA1003433, HEMBA1003538, HEMBA1003555, HEMBA1003568, HEMBA1003569, HEMBA100 3581,
        HEMBA1003720, HEMBA1003760, HEMBA1004168, HEMBA1004248, HEMBA1004276, HEMBA10 04354,
        HEMBA1004356, HEMBA1004479, HEMBA1004534, HEMBA1004669, HEMBA1004752, HEMBA1 004753,
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       HEMBA1004807, HEMBA1005293, HEMBA1005331, HEMBA1005359, HEMBA1005367, HEMBA 1005423,
       HEMBA1005443, HEMBA1005475, HEMBA1005528, HEMBA1005576, HEMBA1005581, HEMB A1005679,
       HEMBA1005699, HEMBA1005732, HEMBA1005931, HEMBA1006173, HEMBA1006474, HEM BA1006562,
       HEMBA1006648, HEMBA1006652, HEMBA1006737, HEMBA1006807, HEMBA1006976, HE MBA1007078,
       HEMBA1007121, HEMBA1007243, HEMBA1007301, HEMBB1000264, HEMBB1000637, H EMBB1000684,
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       HEMBB1000693, HEMBB1000915, HEMBB1000927, HEMBB1001011, HEMBB1001051, HEMBB1001068,
       HEMBB1001102, HEMBB1001119, HEMBB1001133, HEMBB1001177, HEMBB1001802, HEMBB1001908,
       HEMBB1001922, HEMBB1001967, HEMBB1002050, HEMBB1002193, HEMBB100221 7, HEMBB1002358,
       HEMBB1002635, HEMBB1002683, MAMMA1000009, MAMMA1000092, MAMMA10001 03, MAMMA1000173,
       MAMMA1000731, MAMMA1000897, MAMMA1000956, MAMMA1001021,
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       MAMMA1001041, MAMMA1001105, MAMMA1001198, MAMMA1001305, MAMMA1001576, MAMMA1001633,
       MAMMA1002140, MAMMA1002170, MAMMA1002198, MAMMA1002268, MAMMA1002352, MAMMA100239 2,
       MAMMA1002485, MAMMA1002530, MAMMA1002573, MAMMA1002636, MAMMA1002858, MAMMA10028 69,
       MAMMA1002881, MAMMA1002937, MAMMA1003047, MAMMA1003099, MAMMA1003146, MAMMA1003 166,
       NT2RM1000035, NT2RM1000153. NT2RM1000257. NT2RM1000318, NT2RM1000377, NT2RM100 0394,
25
       NT2RM1000555, NT2RM1000691, NT2RM1000725, NT2RM1000770, NT2RM1000826, NT2RM10 00850,
       NT2RM1001059, NT2RM1001072, NT2RM1001082, NT2RM1001092, NT2RM2000363, NT2RM2 000374,
       NT2RM2000594, NT2RM2000599, NT2RM2000714, NT2RM2000740, NT2RM2000821, NT2RM 2001035,
       NT2RM2001499, NT2RM2001575, NT2RM2001605, NT2RM2001664, NT2RM2001670, NT2R M2001803,
       NT2RM2001823, NT2RM2001839, NT2RM4000085, NT2RM4000139, NT2RM4000155, NT2 RM4000290,
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       NT2RM4000689, NT2RM4000895, NT2RM4001320, NT2RM4001566, NT2RM4001629, NT 2RM4001741,
       NT2RM4001819, NT2RM4001823, NT2RM4001865, NT2RM4001940, NT2RM4002093, N T2RM4002146,
       NT2RM4002161, NT2RM4002189, NT2RM4002457, NT2RM4002558, NT2RM4002571, NT2RP1000086,
       NT2RP1000376, NT2RP1000574, NT2RP1000581, NT2RP1000629, NT2RP1000721, NT2RP1000733,
       NT2RP1000738, NT2RP1000825, NT2RP1000833, NT2RP1000860, NT2RP100096 6, NT2RP1001033,
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       NT2RP1001247, NT2RP2000126, NT2RP2000147, NT2RP2000233, NT2RP20002 97, NT2RP2000310,
       NT2RP2000414, NT2RP2000448, NT2RP2000459, NT2RP2000603.
       NT2RP2000812, NT2RP2001127, NT2RP2001327, NT2RP2001440, NT2RP2001520, NT2RP2001663,
       NT2RP2001748, NT2RP2001876, NT2RP2001898, NT2RP2002025, NT2RP2002066, NT2RP200225 9,
       NT2RP2002312, NT2RP2002503, NT2RP2002618, NT2RP2003228, NT2RP2003293, NT2RP20032 95,
40
       NT2RP2003517, NT2RP2003564, NT2RP2003706, NT2RP2003737, NT2RP2003968, NT2RP2004 013,
       NT2RP2004098, NT2RP2004187, NT2RP2004316, NT2RP2004523, NT2RP2004538, NT2RP200 4933,
       NT2RP2005144, NT2RP2005288, NT2RP2005289, NT2RP2005325, NT2RP2005358, NT2RP20 05531,
       NT2RP2005773, NT2RP2006219, NT2RP3000050, NT2RP3000080, NT2RP3000299, NT2RP3 000397,
       NT2RP3000512, NT2RP3000527, NT2RP3001081, NT2RP3001216, NT2RP3001268, NT2RP 3001355,
       NT2RP3001427, NT2RP3001428, NT2RP3001497, NT2RP3001554, NT2RP3001724, NT2R P3001764,
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       NT2RP3001799, NT2RP3001855, NT2RP3002045, NT2RP3002151, NT2RP3002351, NT2 RP3002352,
       NT2RP3002663, NT2RP3002818, NT2RP3002876, NT2RP3002909, NT2RP3002972, NT 2RP3003061,
       NT2RP3003157, NT2RP3003242, NT2RP3003282, NT2RP3003672, NT2RP3004078, N T2RP3004209,
       NT2RP3004399, NT2RP3004424, NT2RP3004475, NT2RP3004480, NT2RP3004490, NT2RP3004527,
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       NT2RP3004578, NT2RP4000049, NT2RP4000109, NT2RP4000185, NT2RP4000312, NT2RP4000367,
       NT2RP4000398, NT2RP4000457, NT2RP4000855, NT2RP4000879, NT2RP400091 5, NT2RP4000928,
       NT2RP4001095, NT2RP4001150, NT2RP4001213, NT2RP4001483, NT2RP40014 98, NT2RP4001753,
       NT2RP4001838, OVARC1000014, OVARC1000114, OVARC1000139,
       OVARC1000241, OVARC1000440, OVARC1000543, OVARC1000722, OVARC1000771, OVARC1000834,
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       OVARC1000850, OVARC1001113, OVARC1001244, OVARC1001296, OVARC1001357, OVARC100137 2,
       OVARC1001417, OVARC1001496, OVARC1001506, OVARC1001577, OVARC1001668, OVARC10017 13,
       OVARC1001809, OVARC1001880, OVARC1001901, OVARC1001949, OVARC1002050, PLACE1000 133,
       PLACE1000142, PLACE1000184, PLACE1000383, PLACE1000406, PLACE1000420, PLACE100 0583,
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PLACE1000588, PLACE1000706, PLACE1001384, PLACE1001387, PLACE1001517, PLACE10 01602,
       PLACE1001634, PLACE1001771, PLACE1002046, PLACE1002437, PLACE1002474, PLACE1 002665,
       PLACE1002834, PLACE1003302, PLACE1003493, PLACE1003704, PLACE1003888, PLACE 1003903,
       PLACE1003968, PLACE1004197, PLACE1004358, PLACE1004471, PLACE1004681, PLAC E1004751,
       PLACE1004773, PLACE1005101, PLACE1006040, PLACE1006119, PLACE1006170, PLA CE1006187,
       PLACE1006288, PLACE1006414, PLACE1006438, PLACE1006962, PLACE1007239, PL ACE1007257,
       PLACE1007488, PLACE1007511, PLACE1007737, PLACE1008000, PLACE1008402, P LACE1008643,
       PLACE1008696, PLACE1008941, PLACE1009027, PLACE1009200, PLACE1009230, PLACE1009298,
       PLACE1009319, PLACE1009444, PLACE1009524, PLACE1010083, PLACE1010089, PLACE1010599,
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       PLACE1010833, PLACE1010856, PLACE1011054, PLACE1011057, PLACE101116 5, PLACE1011229,
       PLACE1011263, PLACE1011371, PLACE2000072, PLACE2000399, PLACE20004 38, PLACE2000458,
       PLACE3000004, PLACE3000242, PLACE3000352, PLACE3000353,
       PLACE3000455, PLACE3000475, PLACE4000009, PLACE4000014, PLACE4000052, PLACE4000320,
       PLACE4000581, SKNMC1000050, THYRO1000026, THYRO1000173, THYRO1000186, THYRO100032 0,
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       THYRO1000327, THYRO1000387, THYRO1000471, THYRO1000934, THYRO1001458, THYRO10015 95,
       THYRO1001605, THYRO1001617, THYRO1001671, Y79AA1000037, Y79AA1000469, Y79AA1000 539,
       Y79AA1000560, Y79AA1000734, Y79AA1000985, Y79AA1001105, Y79AA1001391, Y79AA100 1394,
       Y79AA1001548, Y79AA1001613, Y79AA1001874, Y79AA1002258, Y79AA1002416, Y79AA10 02482,
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20 (6) The following 301 clones were predicted to encode a protein in the category of enzyme or proteins associated with metabolism (including the clones belonging to plural categories). HEMBA1000012, HEMBA1000459, HEMBA1000852, HEMBA1001059, HEMBA1001257, HEMBA1001620, HEMBA1001714, HEMBA1001744, HEMBA1001896, HEMBA1002003, HEMBA1002212, HEMBA100225 7, HEMBA1002513, HEMBA1002973, HEMBA1003046, HEMBA1003136, HEMBA1003179, HEMBA10032 86, 25 HEMBA1003291, HEMBA1003538, HEMBA1003680, HEMBA1004227, HEMBA1004408, HEMBA1004 734, HEMBA1004795, HEMBA1005513, HEMBA1005737, HEMBA1005815, HEMBA1006278, HEMBA100 6347, HEMBA1006521, HEMBA1006648, HEMBA1006877, HEMBA1006941, HEMBA1006976, HEMBA10 07121, HEMBA1007243, HEMBA1007300, HEMBB1000781, HEMBB1000915, HEMBB1001288, HEMBB1 001429, HEMBB1001443, HEMBB1001915, HEMBB1002042, HEMBB1002342, HEMBB1002358, HEMBB 1002635, HEMBB1002683, MAMMA1000020, MAMMA1000085, MAMMA1000672, MAMMA1000841, MAMM A1001008, MAMMA1001139, MAMMA1001476, MAMMA1001501, MAMMA1002268, MAMMA1002530, MAM MA1002619, MAMMA1002671, MAMMA1002938, NT2RM1000132, NT2RM1000153, NT2RM1000256, NT 2RM1000280, NT2RM1000377, NT2RM1000648, NT2RM1000850, NT2RM1000867, NT2RM1000894, N T2RM1001072, NT2RM2000013, NT2RM2000124, NT2RM2000191, NT2RM2000368, NT2RM2000371, NT2RM2000577, 35 NT2RM2000594, NT2RM2000599, NT2RM2000951, NT2RM2001238, NT2RM2001664, NT2RM2001700, NT2RM2001730, NT2RM2001785, NT2RM2001803, NT2RM2002030, NT2RM400002 4, NT2RM4000155, NT2RM4000344, NT2RM4000496. NT2RM4000616, NT2RM4000712, NT2RM40008 95, NT2RM4001313, NT2RM4001316, NT2RM4001340, NT2RM4001444, NT2RM4001758, NT2RM4001819, NT2RM4001930, NT2RM4002062, NT2RM4002063, NT2RM4002409, NT2RM4002558, 40 NT2RM4002571, NT2RM4002623, NT2RP1000018, NT2RP1000112, NT2RP1000376, NT2RP100060 9, NT2RP1000833, NT2RP1000834, NT2RP1001079, NT2RP1001253, NT2RP1001361, NT2RP10015 43, NT2RP2000056, NT2RP2000114, NT2RP2000183, NT2RP2000329, NT2RP2000422, NT2RP2000 710, NT2RP2000764, NT2RP2000932, NT2RP2001070, NT2RP2001663, NT2RP2001839, NT2RP200 1898, NT2RP2002256, NT2RP2002312, NT2RP2002442, NT2RP2002595, NT2RP2002618, NT2RP20 02993, 45 NT2RP2003230, NT2RP2003286, NT2RP2003506, NT2RP2003643, NT2RP2003706, NT2RP2 003737, NT2RP2003781, NT2RP2003912, NT2RP2004232, NT2RP2004239, NT2RP2004768, NT2RP 2004791, NT2RP2004799, NT2RP2005038, NT2RP2005276, NT2RP2005344, NT2RP2005457, NT2R P2005498, NT2RP2005531, NT2RP2005557, NT2RP2005690, NT2RP2005773, NT2RP2005775, NT2 RP2005942, NT2RP3000571, NT2RP3000046, NT2RP3000085, NT2RP3000359, NT2RP3000418, NT 2RP3000742, 50 NT2RP3000845, NT2RP3000875, NT2RP3001274, NT2RP3001407, NT2RP3001495, N T2RP3002351, NT2RP3002402, NT2RP3002969, NT2RP3003301, NT2RP3003346, NT2RP3003500, NT2RP3003800, NT2RP3003825, NT2RP3003831, NT2RP3003914, NT2RP3004155, NT2RP3004209 NT2RP3004669, NT2RP3004670, NT2RP4000259, NT2RP4000417, NT2RP4000457, NT2RP400072 4, NT2RP4000855, NT2RP4000928, NT2RP4000997, NT2RP4001041, NT2RP4001079, NT2RP40010 95, NT2RP4001219, 55 NT2RP4001345, NT2RP4001375, NT2RP4001483, NT2RP4001592, NT2RP4001644, NT2RP4001893, NT2RP4001946, NT2RP4002408, NT2RP5003500, NT2RP5003522, OVARC1000060, OVARC1000139, OVARC1000309, OVARC1000543, OVARC1000682, OVARC100072 2,

OVARC1000885, OVARC1001261, OVARC1001610, OVARC1001713, OVARC1001762, OVARC10018 09,

OVARC1001942, PLACE1000007, PLACE1000142, PLACE1000420, PLACE1000547, PLACE1000 653, PLACE1001062, PLACE1001603, PLACE1001692, PLACE1001817, PLACE1001989, PLACE100 2991, PLACE1003100, PLACE1003709, PLACE1003885, PLACE1003888, PLACE1003903, PLACE10 03915, PLACE1003968, PLACE1004428, PLACE1004437, PLACE1004751, PLACE1005305, PLACE1 005477, PLACE1005656, PLACE1005763, PLACE1005804, PLACE1006011, PLACE1016469, PLACE 1006731, PLACE1007729, PLACE1007958, PLACE1008275, PLACE1008356, PLACE1008696, PLAC E1009444, PLACE1010481, PLACE1011046, PLACE1010089, PLACE1010096, PLACE1010106, PLA CE1010152, PLACE1010481, PLACE1011046, PLACE1011203, PLACE1011219, PLACE1011229, PLACE1011322, PLACE3000148, PLACE3000156, PLACE3000350, PLACE3000404, PLACE2000411, PLACE3000020, PLACE3000148, PLACE3000590, SKNMC1000050, THYRO1000017, THYRO1000197, THYRO1000320, THYRO1000358, THYRO1000471, THYRO1000926, THYRO1000934, THYRO100095 1, THYRO1001406, THYRO1001617, THYRO1001671, Y79AA1001048, Y79AA1001233, Y79AA10015 81, Y79AA1001679, Y79AA1001827, Y79AA1002027, Y79AA1002209, Y79AA1002361, Y79AA1002 416,

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Y79AA1000214, Y79AA1001236,

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- (7) The following 74 clones were predicted to encode a protein in the category of proteins associated with cell division or cell proliferation (including the clones belonging to plural categories).

  HEMBA1001019, HEMBA1001595, HEMBA1002229, HEMBA1002363, HEMBA1003136, HEMBA1003836, HEMBA1004131, HEMBA1005241, HEMBB1000399, HEMBB1001175, HEMBB1001242, MAMMA100182 0, MAMMA1003011, NT2RM1000394, NT2RM2001256, NT2RM4000215, NT2RM4001611, NT2RM40017 14, NT2RP1000733, NT2RP1001177, NT2RP2000077, NT2RP2000346, NT2RP2001397, NT2RP2003 912, NT2RP2005520, NT2RP2005675, NT2RP2005767, NT2RP2005857, NT2RP2006464, NT2RP300 0031, NT2RP3001155, NT2RP3001730, NT2RP3002151, NT2RP3002273, NT2RP3002399, NT2RP30 04507, NT2RP3004578, NT2RP3004594, NT2RP4000210, NT2RP4000498, NT2RP4001207, NT2RP4 001414, NT2RP4001551, OVARC1000066, OVARC1000087, OVARC1000113, OVARC1000846, OVARC 1000876, OVARC1001068, OVARC1001107, OVARC1001113, OVARC1002112, PLACE100547, PLAC E1001611, PLACE1004814, PLACE1005287, PLACE1006187, PLACE1006368, PLACE1008808, PL ACE1008947,
- (8) The following 92 clones were predicted to encode a protein in the category of proteins associated with cytoskeleton (including the clones belonging to plural categories).

PLACE1009060, PLACE1010720, PLACE1010761, PLACE1010833, PLACE1011056, P LACE3000244,

HEMBA1000020, HEMBA1000588, HEMBA1001804, HEMBA1002102, HEMBA1002161, HEMBA1002939, HEMBA1003235, HEMBA1003581, HEMBA1004534, HEMBA1005009, HEMBA1005595, HEMBA100627 2, HEMBA1006737, HEMBA1007018, HEMBB1000404, HEMBB1001282, MAMMA1000173, MAMMA10004 29, MAMMA1001041, MAMMA1001576, MAMMA1001735, MAMMA1002622, MAMMA1002637, MAMMA1003 099, MAMMA1003127, NT2RM1000898, NT2RM1001003, NT2RM1001139, NT2RM2000691, NT2RM400 0030, NT2RM4000167, NT2RM4000689, NT2RM4001741, NT2RM4001776, NT2RM4002109, NT2RP10 00348, NT2RP1000460, NT2RP1000478, NT2RP1001033, NT2RP2000812, NT2RP2001634, NT2RP2 001900, NT2RP2002727, NT2RP2003307, NT2RP2003604, NT2RP2004538, NT2RP2005531, NT2RP 3000868, NT2RP3001216, NT2RP3001272, NT2RP3002688, NT2RP3003061, NT2RP3003138, NT2R P3003230, NT2RP3003282, NT2RP3004454, NT2RP3004569, NT2RP4001126, NT2RP4001927, OVA RC1000437, OVARC1000520, OVARC1000679, OVARC1001731, OVARC1002050, PLACE1002571, PL ACE1002591, PLACE1002655, PLACE1003669, PLACE1004777, PLACE1005287, PLACE1006368, P LACE1007105, PLACE1010310, PLACE1010743, PLACE1010960, PLACE1011263, PLACE2000039, PLACE2000216, PLACE2000274, PLACE3000145, PLACE3000322, PLACE4000009, PLACE4000612, THYRO1000072, THYRO1000666, THYRO1001121, THYRO1001458, Y79AA1000368, Y79AA100083 3, Y79AA1000962, Y79AA1000985, Y79AA1002208,

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(9) The following 280 clones were predicted to encode a protein in the category of proteins associated with RNA synthesis (including the clones belonging to plural categories).

HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000591, HEMBA1000851, HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA100181 9, HEMBA1001847, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA10034 18, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004 097,

HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA1004669, HEMBA1004758, HEMBA100 5359, HEMBA1005528, HEMBA1005548, HEMBA1005931, HEMBA1006253, HEMBA1006278, HEMBA10 06359,

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HEMBA1006559, HEMBA1006639, HEMBB1000250, HEMBB1000789, HEMBB1001011, HEMBB1 001314,
        HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAMMA 1000183,
        MAMMA1000388, MAMMA1001059, MAMMA1001633, MAMMA1001820, MAMMA1001837, MAMM A1002650,
        MAMMA1002937, NT2RM1000055, NT2RM1000187, NT2RM1000666, NT2RM1000691, NT2 RM1000852,
        NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT 2RM2001424,
        NT2RM2001575, NT2RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, N T2RM2001989,
        NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000104, NT2RM4000191, NT2RM4000421,
        NT2RM4000471, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996,
        NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM400174 6, NT2RM4001783,
10
        NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RM40020 93, NT2RM4002479,
        NT2RP1000111, NT2RP1000272, NT2RP1000333, NT2RP1000470,
        NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1001080, NT2RP2000008, NT2RP2000133,
        NT2RP2000161, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP200144 9,
        NT2RP2002099, NT2RP2002252, NT2RP2002503, NT2RP2002591, NT2RP2002928, NT2RP20031 25,
15
        NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004 187,
        NT2RP2004568, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005126, NT2RP200 5168,
        NT2RP2005239, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP20 05763,
        NT2RP2005942, NT2RP2006238, NT2RP2006464, NT2RP3000050, NT2RP3000361, NT2RP3 000397,
        NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000632, NT2RP3000917, NT2RP 3001057,
20
       NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001399, NT2R P3001671,
        NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP3001855, NT2 RP3002165,
        NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP3003251, NT 2RP3003327,
        NT2RP3004013, NT2RP3004258, NT2RP3004504, NT2RP3004566, NT2RP3004572, N T2RP3004594,
       NT2RP3004617, NT2RP400078, NT2RP4000111, NT2RP4000210, NT2RP4000398, NT2RP4000449,
25
       NT2RP4000518, NT2RP4000614, NT2RP4000787, NT2RP4000865, NT2RP4000997 . NT2RP4001029,
       NT2RP4001080, NT2RP4001148, NT2RP4001213, NT2RP4001433, NT2RP400152 9, NT2RP4001551,
       NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP40020 78, OVARC1000479,
       OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577,
       OVARC1001949, PLACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706,
30
       PLACE1001118, PLACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE100245 0,
       PLACE1002532, PLACE1002834, PLACE1003190, PLACE1003302, PLACE1003605, PLACE10037 04,
       PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004 681,
       PLACE1004902, PLACE1005646, PLACE1005876, PLACE1006196, PLACE1006414, PLACE100 6438,
       PLACE1006482, PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10 07955,
       PLACE1007969, PLACE1008627, PLACE1008941, PLACE1009099, PLACE1009246, PLACE1 009398,
35
       PLACE1010053, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010580, PLACE 1010702,
       PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072, PLACE2000359, PLAC E4000128,
       PLACE4000156, PLACE4000259, PLACE4000431, THYRO1000121, THYRO1000242, THY RO1000501,
       THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000349, Y7 9AA1000539,
       Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1001105, Y79AA1001299, Y 79AA1001391,
40
       Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1001963, Y79AA1002093,
       Y79AA1002472, Y79AA1002482.
       (10) The following 352 clones were predicted to encode a protein in the category of nuclear protein (including the
45
       clones belonging to plural categories).
       HEMBA1000005, HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000827,
       HEMBA1001851, HEMBA1001088, HEMBA1001137, HEMBA1001476, HEMBA1001510, HEMBA100157 9,
       HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1002035, HEMBA10020 92,
       HEMBA1002363, HEMBA1002495, HEMBA1002569, HEMBA1002609, HEMBA1002999, HEMBA1003 148,
50
       HEMBA1003418, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA100 3783,
       HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA10 04669,
       HEMBA1004758, HEMBA1005359, HEMBA1005475, HEMBA1005528, HEMBA1005548, HEMBA1 005931,
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HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006474, HEMBA1006559, HEMBA 1006914, HEMBB1000250, HEMBB1000399, HEMBB1000789, HEMBB1001011, HEMBB1001314, HEMB B1001482, HEMBB1001908, HEMBB1001915, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAM MA1000183, MAMMA1000388, MAMMA1000731, MAMMA1001633, MAMMA1001820, MAMMA1001837, MA MMA1002650, MAMMA1002869, MAMMA1002937, MAMMA1003011, NT2RM1000055, NT2RM1000187, N T2RM1000257, NT2RM1000394, NT2RM1000666, NT2RM1000691, NT2RM1000852, NT2RM1000894, NT2RM1001092,

NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2RM2001605, NT2RM2001635,

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NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2001823, NT2RM200193 6, NT2RM2001989,
        NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000085, NT2RM40001 04, NT2RM4000191,
        NT2RM4000215, NT2RM4000290, NT2RM4000421, NT2RM4000496,
        NT2RM4000531. NT2RM4000734. NT2RM4000751. NT2RM4000996, NT2RM4001140, NT2RM4001200,
        NT2RM4001217, NT2RM4001382, NT2RM4001483, NT2RM4001592, NT2RM4001605, NT2RM400161 1,
        NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM40019 79,
        NT2RM4002093, NT2RM4002146, NT2RM4002527, NT2RP1000035, NT2RP1000111, NT2RP1000 333,
        NT2RP1000493, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1000966, NT2RP100 1177,
10
        NT2RP2000008, NT2RP2000126, NT2RP2000133, NT2RP2000161, NT2RP2000297, NT2RP20 00414,
        NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2001420, NT2RP2001449, NT2RP2 001536,
        NT2RP2001762, NT2RP2002185, NT2RP2002252, NT2RP2002259, NT2RP2002503, NT2RP 2002504,
        NT2RP2002591, NT2RP2002993, NT2RP2003125, NT2RP2003228, NT2RP2003277, NT2R P2003286,
        NT2RP2003293, NT2RP2003308, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2 RP2004187,
15
       NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT 2RP2005520,
       NT2RP2005539, NT2RP2005722, NT2RP2005767, NT2RP2005857, NT2RP2005942, N T2RP2006238,
       NT2RP3006464, NT2RP3000031, NT2RP3000050, NT2RP3000361, NT2RP3000397, NT2RP3000512,
       NT2RP3000527, NT2RP1000590, NT2RP3000603, NT2RP3000632, NT2RP3000917, NT2RP3001057,
       NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001274, NT2RP300139 8, NT2RP3001399,
       NT2RP3001427, NT2RP3001428. NT2RP3001587, NT2RP3001671, NT2RP30016 72, NT2RP3001688,
20
       NT2RP3001712, NT2RP3001724, NT2RP3001792, NT2RP3001855,
       NT2RP3002056, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193,
       NT2RP3003212, NT2RP3003251, NT2RP3003327, NT2RP3004206, NT2RP3004209, NT2RP300425 8,
       NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3004617, NT2RP4000078, NT2RP40001 11,
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       NT2RP4000150, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000457, NT2RP4000 518,
       NT2RP4000787, NT2RP4000865, NT2RP4000997, NT2RP4001029, NT2RP4001080, NT2RP400 1148,
       NT2RP4001207, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2RP4001551, NT2RP40 01638,
       NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVARC1000006, OVARC1 000087,
       OVARC1000139, OVARC1000326, OVARC1000440, OVARC1000479, OVARC1000846, OVARC 1000883,
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       OVARC1001180, OVARC1001244, OVARC1001381, OVARC1001555, OVARC1001577, OVAR C1001702,
       OVARC1001949, OVARC1002112, PLACE100066, PLACE1000133, PLACE1000406, PLA CE1000583,
       PLACE1000596, PLACE1000706, PLACE1001118, PLACE1001238, PLACE1001304, PL ACE1001602,
       PLACE1001611, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, P LACE1003100,
       PLACE1003190, PLACE1003302, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885,
35
       PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004681, PLACE1004902, PLACE1005287,
       PLACE1005876, PLACE1006011, PLACE1006119, PLACE1006414, PLACE100643 8, PLACE1006482,
       PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10079 55, PLACE1007969,
       PLACE1008044, PLACE1008177, PLACE1008603, PLACE1008627,
       PLACE1008941, PLACE1008947, PLACE1009099, PLACE1009113, PLACE1009246, PLACE1009398,
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       PLACE1010152, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010702, PLACE101072 0,
       PLACE1010771, PLACE1010870, PLACE1011056, PLACE1011229, PLACE1011576, PLACE10116 64,
       PLACE2000072, PLACE2000359, PLACE2000411, PLACE4000014, PLACE4000128, PLACE4000 156,
       PLACE4000259, PLACE4000261, PLACE4000431, THYRO1000121, THYRO1000242, THYRO100 0501,
       THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000214, Y79AA10 00539,
45
       Y79AA1000589, Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1000784, Y79AA1 001105,
       Y79AA1001299, Y79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA 1001866,
       Y79AA1001963, Y79AA1002093, Y79AA1002472, Y79AA1002482,
       (11) The following 111 clones were predicted to encode a protein in the category of proteins associated with protein
50
       synthesis or transport (including the clones belonging to plural categories).
       HEMBA1000012, HEMBA1000542, HEMBA1001913, HEMBA1003773, HEMBA1004202, HEMBA1004356,
       HEMBA1004847, HEMBA1005047, HEMBA1005202, HEMBA1005963, HEMBA1006310, HEMBA100665 2,
       HEMBA1006914, HEMBB1000217, HEMBB1000725, HEMBB1001346, HEMBB1001831, MAMMA10000 85,
       MAMMA1000734, MAMMA1002170, MAMMA1002236, MAMMA1002598, NT2RM1000661, NT2RM2000 504,
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NT2RM2000577, NT2RM2001131, NT2RM4000155, NT2RM4001444, NT2RM4002062, NT2RM400 2205, NT2RM4002623, NT2RP1000326, NT2RP1000947, NT2RP1001569, NT2RP2000710, NT2RP20 00880, NT2RP2001290, NT2RP2001511, NT2RP2001660, NT2RP2002185, NT2RP2002606, NT2RP2 002959, NT2RP2002980, NT2RP2003158, NT2RP2003760, NT2RP2004194, NT2RP2004791, NT2RP 2005012,

NT2RP2005116, NT2RP2005204, NT2RP2006565, NT2RP2006598, NT2RP3000047, NT2R P3000366, NT2RP3000759, NT2RP3000968, NT2RP3001587, NT2RP3002529, NT2RP3003385, NT2 RP3003589, NT2RP3003876, NT2RP3004618, NT2RP4000370, NT2RP4000524, NT2RP4000528, NT 2RP4001041, NT2RP4001574, NT2RP4001592, NT2RP4002047, OVARC1000004, OVARC1000771, O VARC1000862, OVARC1001180, OVARC1001342, OVARC1001766, OVARC1002138, PLACE1000061, PLACE1000492, PLACE1000863, PLACE1001748, PLACE1002090, PLACE1003394, PLACE1003915, PLACE1004149, PLACE1004743, PLACE1005557, PLACE1005813, PLACE1006488, PLACE100653 1, PLACE1006615, PLACE1007706, PLACE1008273, PLACE1008790, PLACE1008813, PLACE10097 21, PLACE1009763, PLACE1009845, PLACE1010074, PLACE1011109, PLACE1011477,

PLACE2000404, PLACE3000059, PLACE3000121, PLACE4000558, PLACE4000654, SKNMC1000011, THYRO1000288, THYRO1000974, Y79AA1000346, Y79AA1000968, Y79AA1002209,

- (12) The following 23 clones were predicted to encode a protein in the category of proteins associated with cellular defense (including the clones belonging to plural categories).
- HEMBA1000005, HEMBA1000531, HEMBB1000217, HEMBB1000399, MAMMA1000734, NT2RP1000493, NT2RP1000493, NT2RP2000006, NT2RP2000045, NT2RP2001536, NT2RP2005204, NT2RP300059 0, NT2RP3001426, NT2RP3002056, NT2RP3004262, NT2RP4001638, OVARC1001900, PLACE10056 11, PLACE1006958, PLACE1008275, PLACE1009113, PLACE4000014, Y79AA1002139.
- (13) The following 23 clones were predicted to encode a protein in the category of proteins associated with development or growth (including the clones belonging to plural categories).
   HEMBB1001802, HEMBB1002442, NT2RM2002142, NT2RM4001047, NT2RP2003308, NT2RP2004816, NT2RP3000994, NT2RP3001340, NT2RP3004206, NT2RP3004472, NT2RP4000246, NT2RP400100 6, OVARC1000304, OVARC1000746, OVARC1000996, PLACE1006037, PLACE1009622, PLACE10116 64, Y79AA1001692.

Table 9
Selected clones having the maximal ATGpr1 score of 0.3 or higher (3690 clones).

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	clone	name of	maximal
	name	sequence	ATGpr1
35			score
	HEMBA1000005	F-HEMBA100005	0. 84
	HEMBA1000012	F-HEMBA1000012	0. 56
40	HEMBA1000020	F-HEMBA1000020	0. 94
40	HEMBA1000030	F-HEMBA1000030	0. 44
	HEMBA1000046	F-HEMBA1000046	0.50
	HEMBA1000050	F-HEMBA1000050	0.94
45	HEMBA1000076	F-HEMBA1000076	0. 48
	HEMBA1000129	F-HEMBA1000129	0.74
	HEMBA1000141	F-HEMBA1000141	0.55
	HEMBA1000150	F-HEMBA1000150	0. 72
50	HEMBA1000156	F-HEMBA1000156	0.94
	HEMBA1000158	F-HEMBA1000158	0. 62
	HEMBA1000168	F-HEMBA1000168	0. 94
	HEMBA1000185	F-HEMBA1000185	0.86
55	HEMBA1000193	F-HEMBA1000193	0.94

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	UEMPA1000001	E 11511011000001	
	HEMBA1000201	F-HEMBA1000201	0.83
	HEMBA1000216	F-HEMBA1000216	0. 32
5	HEMBA1000227	F-HEMBA1000227	0. 72
5	HEMBA1000231	F-HEMBA1000231_	0, 34
	HEMBA1000244	F-HEMBA1000244	0. 51
	HEMBA1000288	F-HEMBA1000288	0.36
	HEMBA1000290	F-HEMBA1000290	0. 45
10	HEMBA1000302	F-HEMBA1000302	0.71
	HEMBA1000303	F-HEMBA1000303	0.42
	HEMBA1000304	F-HEMBA1000304	0. 45
	HEMBA1000307	F-HEMBA1000307	0.48
15	HEMBA1000327	F-HEMBA1000327	0.94
	HEMBA1000356	F-HEMBA1000356	0.90
	HEMBA1000369	F-HEMBA1000369,	0. 94
	HEMBA1000387	F-HEMBA1000387	0.66
20	HEMBA1000392	F-HEMBA1000392	0.60
	HEMBA1000396	F-HEMBA1000396	0.81
	HEMBA1000456	F-HEMBA1000456	0. 90
	HEMBA1000460	F-HEMBA1000460	0. 31
25	HEMBA1000488	F-HEMBA1000488	0.47
25	HEMBA1000490	F-HEMBA1000490	0.89
	HEMBA1000491	F-HEMBA1000491	0.94
	HEMBA1000501	F-HEMBA1000501	0. 67
	HEMBA1000505	F-HEMBA1000505	0.65
30	HEMBA1000508	F-HEMBA1000508	0.39
	HEMBA1000520	F-HEMBA1000520	0. 94
	HEMBA1000523	F-HEMBA1000523	0.86
	HEMBA1000531	F-HEMBA1000531	0. 49
35	HEMBA1000534	F-HEMBA1000534	0. 52
	HEMBA1000542	F-HEMBA1000542	0.81
	HEMBA1000555	F-HEMBA1000555	0. 67
	HEMBA1000561	F-HEMBA1000561	0.34
40	HEMBA1000568	F-HEMBA1000568	0.66
	HEMBA1000588	F-HEMBA1000588	0. 79
	HEMBA1000591	F-HEMBA1000591	0. 92
	HEMBA1000592	F-HEMBA1000592	0.73
45	HEMBA1000594	F-HEMBA1000594	0.89
40	HEMBA1000608	F-HEMBA1000608	0.94
	HEMBA1000636	F-HEMBA1000636	0.94
	HEMBA1000637	F-HEMBA1000637	0. 92
	HEMBA1000657	F-HEMBA1000657	0.85
50	HEMBA1000682	F-HEMBA1000682	0.94
	HEMBA1000686	F-HEMBA1000686	0.90
	HEMBA1000719	F-HEMBA1000719	0. 94
	HEMBA1000727	F-HEMBA1000727	0.40
55	HEMBA1000752	F-HEMBA1000752	0. 31
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	HEMBA1000817	F-HEMBA1000817	0.94
	HEMBA1000851	F-HEMBA1000851	0. 63
_	HEMBA1000867	F-HEMBA1000867	0.68
5	HEMBA1000869	F-HEMBA1000869	0.82
	HEMBA1000872	F-HEMBA1000872	0.54
	HEMBA1000910	F-HEMBA1000910	0.91
	HEMBA1000918	F-HEMBA1000918	0. 38
10	HEMBA1000919	F-HEMBA1000919	0.83
	HEMBA1000942	F-HEMBA1000942	0.31
	HEMBA1000946	F-HEMBA1000946	0.75
	HEMBA1000968	F-HEMBA1000968	0.39
15	HEMBA1000971	F-HEMBA1000971	0.74
	HEMBA1000975	F-HEMBA1000975	0.46
	HEMBA1000986	F-HEMBA1000986,	0.89
	HEMBA1001008	F-HEMBA1001008	0.47
20	HEMBA1001009	F-HEMBA1001009	0.86
	HEMBA1001022	F-HEMBA1001022	0.57
	HEMBA1001043	F-HEMBA1001043	0. 79
	HEMBA1001052	F-HEMBA1001052	0.83
25	HEMBA1001059	F-HEMBA1001059	0.78
	HEMBA1001080	F-HEMBA1001080	0.77
	HEMBA1001085	F-HEMBA1001085	0.92
	HEMBA1001088	F-HEMBA1001088	0.94
30	HEMBA1001109	F-HEMBA1001109	0.50
30	HEMBA1001122	F-HEMBA1001122	0.74
	HEMBA1001133	F-HEMBA1001133	0.45
	HEMBA1001137	F-HEMBA1001137	0.37
25	HEMBA1001140	F-HEMBA1001140	0.43
35	HEMBA1001174	F-HEMBA1001174	0. 44
	HEMBA1001197	F-HEMBA1001197	0.65
	HEMBA1001213	F-HEMBA1001213	0. 61
	HEMBA1001235	F-HEMBA1001235	0.94
40	HEMBA1001247	F-HEMBA1001247	0.31
	HEMBA1001257	F-HEMBA1001257	0.92
	HEMBA1001281	F-HEMBA1001281	0.94
	HEMBA1001286	F-HEMBA1001286	0.49
45	HEMBA1001289	F-HEMBA1001289	0.94
	HEMBA1001302	F-HEMBA1001302	0.62
	HEMBA1001303	F-HEMBA1001303	0.79
	HEMBA1001310	F-HEMBA1001310	0. 66
50	HEMBA1001326	F-HEMBA1001326	0.59
	HEMBA1001351	F-HEMBA1001351	0. 65
	HEMBA1001361	F-HEMBA1001361	0.94
	HEMBA1001377	F-HEMBA1001377	0.87
55	HEMBA1001387	F-HEMBA1001387	0. 94
	HEMBA1001388	F-HEMBA1001388	0. 57

	HEMBA1001398	F-HEMBA1001398	0.74
	HEMBA1001405	F-HEMBA1001405	0.61
	HEMBA1001407	F-HEMBA1001407	0.43
5	HEMBA1001413	F-HEMBA1001413	0.40
	HEMBA1001415	F-HEMBA1001415	0. 94
	HEMBA1001446	F-HEMBA1001446	0. 90
	HEMBA1001450	F-HEMBA1001450	0.40
10	HEMBA1001455	F-HEMBA1001455	0. 73
	HEMBA1001476	F-HEMBA1001476	0. 65
	HEMBA1001497	F-HEMBA1001497	0.48
	HEMBA1001510	F-HEMBA1001510	0.36
15	HEMBA1001516	F-HEMBA1001516	0. 66
	HEMBA1001533	F-HEMBA1001533	0. 46
	HEMBA1001569 HEMBA1001570	F-HEMBA1001569,	
	HEMBA1001570	F-HEMBA1001570 F-HEMBA1001579	0. 53 0. 82
20			
	HEMBA1001581	F-HEMBA1001581	0.80
	HEMBA1001595	F-HEMBA1001595	0.72
	HEMBA1001620	F-HEMBA1001620	0.94
25	HEMBA1001635	F-HEMBA1001635	0.87
	HEMBA1001640	F-HEMBA1001640	0.82
	HEMBA1001647	F-HEMBA1001647	0.91
	HEMBA1001655	F-HEMBA1001655	0.39
30	HEMBA1001661	F-HEMBA1001661	0.86
	HEMBA1001672	F-HEMBA1001672	0. 94
	HEMBA1001702	F-HEMBA1001702	0.72
	HEMBA1001711	F-HEMBA1001711	0. 74
35	HEMBA1001714	F-HEMBA1001714	0.89
	HEMBA1001723	F-HEMBA1001723	0.94
	HEMBA1001731	F-HEMBA1001731	0.37
	HEMBA1001744	F-HEMBA1001744	0.31
	HEMBA1001746	F-HEMBA1001746	0. 39
40	HEMBA1001781	F-HEMBA1001781	0.67
	HEMBA1001800	F-HEMBA1001800	0. 69
	HEMBA1001804		
	HEMBA1001809	F-HEMBA1001809	0. 78
45	HEMBA1001815	F-HEMBA1001815	0.49
	HEMBA1001819	F-HEMBA1001819	0. 90
	HEMBA1001822	F-HEMBA1001822	0.80
	HEMBA1001824	F-HEMBA1001824	0. 68
50	HEMBA1001847	F-HEMBA1001847	0.46
	HEMBA1001864	F-HEMBA1001864	0.34
	HEMBA1001866	F-HEMBA1001866	0. 53
	HEMBA1001869	F-HEMBA1001869	0.69
55	HEMBA1001896	F-HEMBA1001896	0. 48
	HEMBA1001910	F-HEMBA1001910	0.84

	HEMBA1001912	F-HEMBA1001912	0.48
	HEMBA1001913	F-HEMBA1001913	0.94
	HEMBA1001915	F-HEMBA1001915	0.41
5	HEMBA1001921	F-HEMBA1001921	0. 94
	HEMBA1001939	F-HEMBA1001939	0.87
	HEMBA1001950	F-HEMBA1001950	0. 36
	HEMBA1001967	F-HEMBA1001967	0.94
10	HEMBA1001987	F-HEMBA1001987	0.41
	HEMBA1002018	F-HEMBA1002018	0. 41
	HEMBA1002035	F-HEMBA1002035	0.36
	HEMBA1002049	F-HEMBA1002049	0.39
15	HEMBA1002084	F-HEMBA1002084	0. 45
	HEMBA1002092	F-HEMBA1002092	0. 94
	HEMBA1002102	F-HEMBA1002102,	0. 94
	HEMBA1002119	F-HEMBA1002119	0.60
20	HEMBA1002125	F-HEMBA1002125	0.88
	HEMBA1002150	F-HEMBA1002150	0. 31
	HEMBA1002151	F-HEMBA1002151	0.82
	HEMBA1002161	F-HEMBA1002161	0.94
25	HEMBA1002177	F-HEMBA1002177	0.38
	HEMBA1002189	F-HEMBA1002189	0. 43
	HEMBA1002191	F-HEMBA1002191	0.94
	HEMBA1002199	F-HEMBA1002199	0. 85
30	HEMBA1002212	F-HEMBA1002212	0. 39
	HEMBA1002215	F-HEMBA1002215	0. 94
	HEMBA1002229	F-HEMBA1002229	0.83
	HEMBA1002237	F-HEMBA1002237	0. 86
35	HEMBA1002241	F-HEMBA1002241	0.66
33	HEMBA1002265	F-HEMBA1002265	0. 38
	HEMBA1002267	F-HEMBA1002267	0. 33
	HEMBA1002341	F-HEMBA1002341	0. 53
-	HEMBA1002363	F-HEMBA1002363	0.88
40	HEMBA1002417	F-HEMBA1002417	0. 83
	HEMBA1002419	F-HEMBA1002419	0. 92
	HEMBA1002430	F-HEMBA1002430	0. 45
	HEMBA1002439	F-HEMBA1002439	0.83
45	HEMBA1002458	F-HEMBA1002458	0. 47
	HEMBA1002460	F-HEMBA1002460	0. 48
	HEMBA1002462	F-HEMBA1002462	0.55
	HEMBA1002469	F-HEMBA1002469 F-HEMBA1002475	0. 79
50	HEMBA1002475		0. 88 0. 45
	HEMBA1002477	F-HEMBA1002477 F-HEMBA1002495	0. 45
	HEMBA1002495	F-HEMBA1002503	0. 78
	HEMBA1002503 HEMBA1002508	F-HEMBA1002508	0. 94
55		F-HEMBA1002513	0. 33
	HEMBA1002513	F-TIEMDA (UUZD13	0, 41

	HEMBA1002515	F-HEMBA1002515	0.94
	HEMBA1002542		0. 46
	HEMBA1002547		0. 45
5	HEMBA1002569	F-HEMBA1002569	
	HEMBA1002583	F-HEMBA1002583	0. 86
	HEMBA1002609	F-HEMBA1002609	0. 75
	HEMBA1002624	F-HEMBA1002624	0. 59
10	HEMBA1002688	F-HEMBA1002688	0.88
	HEMBA1002696		0. 82
	HEMBA1002703		0. 94
	HEMBA1002746	F-HEMBA1002746	0. 33
15	HEMBA1002750	F-HEMBA1002750	0. 53
	HEMBA1002768		0. 94
	HEMBA1002770		
	HEMBA1002777		0. 90
20	HEMBA1002779		0. 92
	HEMBA1002794		0.86
	HEMBA1002810		0.87
	HEMBA1002816	F-HEMBA1002816	0. 91
0.5	HEMBA1002818	F-HEMBA1002818	0. 94
25	HEMBA1002850	F-HEMBA1002850	0. 57
	HEMBA1002863	F-HEMBA1002863	0. 48
	HEMBA1002876		0. 86
	HEMBA1002935	F-HEMBA1002935	0. 91
30	HEMBA1002937		0. 61
	HEMBA1002939		0. 32
	HEMBA1002951		0.83
		F-HEMBA1002954	0. 67
35		F-HEMBA1002970	0. 33
	HEMBA1002971	F-HEMBA1002971	0. 57
		F-HEMBA1002973	0. 33
	HEMBA1002997	F-HEMBA1002997	0. 46
40	HEMBA1002999	F-HEMBA1002999	0. 49
	HEMBA1003021	F-HEMBA1003021	0. 74
	HEMBA1003033		0. 94
		F-HEMBA1003035	0. 61
45	HEMBA1003041	F-HEMBA1003041	0. 40
	HEMBA1003046	F-HEMBA1003046	0. 94
	HEMBA1003067	F-HEMBA1003067	0. 34
	HEMBA1003077	F-HEMBA1003077	0. 83
50	HEMBA1003078	F-HEMBA1003078	0. 89
50	HEMBA1003079	F-HEMBA1003079	0. 49
	HEMBA1003096	F-HEMBA1003096	0. 34
	HEMBA1003090	F-HEMBA1003117	0. 94
	HEMBA1003129	F-HEMBA1003129	0. 94
55	HEMBA1003136	F-HEMBA1003136	0. 77
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	HEMBA1003148	F-HEMBA1003148	0.91
	HEMBA1003175	F-HEMBA1003175	0. 41
5	HEMBA1003179	F-HEMBA1003179	0.42
3	HEMBA1003199	F-HEMBA1003199	0.84
	HEMBA1003222	F-HEMBA1003222	0.47
	HEMBA1003235	F-HEMBA1003235	0.94
	HEMBA1003250	F-HEMBA1003250	0.39
10	HEMBA1003257	F-HEMBA1003257	0.88
	HEMBA1003281	F-HEMBA1003281	0. 67
	HEMBA1003286	F-HEMBA1003286	0.57
	HEMBA1003291	F-HEMBA1003291	0.89
15	HEMBA1003322	F-HEMBA1003322	0.60
	HEMBA1003327	F-HEMBA1003327	0. 88
	HEMBA1003369	F-HEMBA1003369,	
	HEMBA1003370	F-HEMBA1003370	0, 75
20	HEMBA1003380	F-HEMBA1003380	0.68
	HEMBA1003395	F-HEMBA1003395	0. 88
	HEMBA1003402	F-HEMBA1003402	0. 45
	HEMBA1003403	F-HEMBA1003403	0.44
25	HEMBA1003408	F-HEMBA1003408	0.34
	HEMBA1003417	F-HEMBA1003417	0.78
	HEMBA1003418	F-HEMBA1003418	0.94
	HEMBA1003433	F-HEMBA1003433	0.70
30	HEMBA1003447	F-HEMBA1003447	0. 49
	HEMBA1003461	F-HEMBA1003461	0.85
	HEMBA1003463	F-HEMBA1003463	0. 59
	HEMBA1003480	F-HEMBA1003480	0.94
35	HEMBA1003528	F-HEMBA1003528	0.90
33	HEMBA1003538	F-HEMBA1003538	0. 47
	HEMBA1003545	F-HEMBA1003545	0. 74
	HEMBA1003555	F-HEMBA1003555	0. 94
40	HEMBA1003556	F-HEMBA1003556	0. 44
40	HEMBA1003560	F-HEMBA1003560	0. 40
	HEMBA1003568	F-HEMBA1003568	0. 94
	HEMBA1003569	F-HEMBA1003569	0. 82
	HEMBA1003581	F-HEMBA1003581	0. 83
45	HEMBA1003591	F-HEMBA1003591	0.68
	HEMBA1003615	F-HEMBA1003615	0.94
	HEMBA1003617	F-HEMBA1003617	0. 88
	HEMBA1003621	F-HEMBA1003621	0. 47
50	HEMBA1003645	F-HEMBA1003645	0. 75
	HEMBA1003646	F-HEMBA1003646	0.51
	HEMBA1003662	F-HEMBA1003662	0. 61
	HEMBA1003667	F-HEMBA1003667	0. 54
55	HEMBA1003679	F-HEMBA1003679	0. 72
	HEMBA1003680	F-HEMBA1003680	0. 52

	HEMBA1003684	F-HEMBA1003684	0.94
	HEMBA1003690	F-HEMBA1003690	0.93
	HEMBA1003692	F-HEMBA1003692	0.46
5	HEMBA1003711	F-HEMBA1003711	0. 94
	HEMBA1003720	F-HEMBA1003720	0. 94
	HEMBA1003729	F-HEMBA1003729	0. 73
	HEMBA1003742	F-HEMBA1003742	0. 94
10	HEMBA1003760	F-HEMBA1003760	0. 75
	HEMBA1003783	F-HEMBA1003783	0. 31
	HEMBA1003799	F-HEMBA1003799	0.70
	HEMBA1003803	F-HEMBA1003803	0.87
15	HEMBA1003804	F-HEMBA1003804	0. 94
	HEMBA1003805	F-HEMBA1003805	0. 94
	HEMBA1003807	F-HEMBA1003807,	0. 91
	HEMBA1003827	F-HEMBA1003827	0.62
20	HEMBA1003836	F-HEMBA1003836	0.69
	HEMBA1003838	F-HEMBA1003838	0.80
	HEMBA1003864	F-HEMBA1003864	0.94
	HEMBA1003866	F-HEMBA1003866	0.32
25	HEMBA1003879	F-HEMBA1003879	0.82
	HEMBA1003880	F-HEMBA1003880	0. 59
	HEMBA1003893	F-HEMBA1003893	0. 47
	HEMBA1003953	F-HEMBA1003953	0. 77
30	HEMBA1003959	F-HEMBA1003959	0. 32
	HEMBA1003978	F~HEMBA1003978	0. 37
	HEMBA1003985	F-HEMBA1003985	0. 31
	HEMBA1003989	F-HEMBA1003989	0. 75
35	HEMBA1004011	F-HEMBA1004011	0. 34
	HEMBA1004048	F-HEMBA1004048	0. 32
	HEMBA1004055	F-HEMBA1004055 F-HEMBA1004056	0. 74
	HEMBA1004056 HEMBA1004074	F-HEMBA1004074	0. 49 0. 39
40	HEMBA1004074	F-HEMBA1004086	0. 35
	HEMBA1004087	F-HEMBA1004097	0. 43
	HEMBA1004111	F-HEMBA1004111	0. 94
	HEMBA1004131	F-HEMBA1004131	0. 84
45	HEMBA1004133	F-HEMBA1004133	0. 40
	HEMBA1004143	F-HEMBA1004143	0. 33
	HEMBA1004146	F-HEMBA1004146	0. 31
	HEMBA1004150	F-HEMBA1004150	0. 70
50	HEMBA1004168	F-HEMBA1004168	0. 34
	HEMBA1004199	F-HEMBA1004199	0.34
	HEMBA1004200	F-HEMBA1004200	0.40
	HEMBA1004202	F-HEMBA1004202	0.88
55	HEMBA1004203	F-HEMBA1004203	0.94
55	HEMBA1004207	F-HEMBA1004207	0.53

	HEMBA1004227	F-HEMBA1004227	0. 94
	HEMBA1004238	F-HEMBA1004238	0. 44
5	HEMBA1004246	F-HEMBA1004246	0.71
3	HEMBA1004248	F-HEMBA1004248	0.40
	HEMBA1004264	F-HEMBA1004264	0. 94
	HEMBA1004274	F-HEMBA1004274	0.88
	HEMBA1004275	F-HEMBA1004275	0, 85
10	HEMBA1004276	F-HEMBA1004276	0. 32
	HEMBA1004286	F-HEMBA1004286	0.37
	HEMBA1004289	F-HEMBA1004289	0.76
	HEMBA1004321	F-HEMBA1004321	0. 82
15	HEMBA1004327	F-HEMBA1004327	0.40
	HEMBA1004330	F-HEMBA1004330	0.84
	HEMBA1004335	F-HEMBA1004335,	0. 79
	HEMBA1004341	F-HEMBA1004341	0.38
20	HEMBA1004353	F-HEMBA1004353	0.79
	HEMBA1004372	F-HEMBA1004372	0.94
	HEMBA1004389	F-HEMBA1004389	0.32
	HEMBA1004408	F-HEMBA1004408	0.70
25	HEMBA1004479	F-HEMBA1004479	0. 55
	HEMBA1004499	F-HEMBA1004499	0. 62
	HEMBA1004502	F-HEMBA1004502	0. 31
	HEMBA1004507	F-HEMBA1004507	0. 71
30	HEMBA1004509	F-HEMBA1004509	0. 68
	HEMBA1004534	F-HEMBA1004534	0. 56
	HEMBA1004542	F-HEMBA1004542	0. 36
	HEMBA1004554	F-HEMBA1004554	0. 90
35	HEMBA1004560	F-HEMBA1004560	0. 57
	HEMBA1004573	F-HEMBA1004573	0. 93
	HEMBA1004596	F-HEMBA1004596	0. 45
	HEMBA1004604	F-HEMBA1004604	0. 91
40	HEMBA1004610	F-HEMBA1004610	0. 59
	HEMBA1004632	F-HEMBA1004632	0. 59
	HEMBA1004637 HEMBA1004638	F-HEMBA1004637	0. 38
		F-HEMBA1004638	0. 40
45	HEMBA1004669 HEMBA1004693	F-HEMBA1004669	0. 94
	HEMBA1004697	F-HEMBA1004693	0. 52
	HEMBA1004705	F-HEMBA1004697 F-HEMBA1004705	0.94
	HEMBA1004709	F-HEMBA1004709	0.94
50	HEMBA1004709	F-HEMBA1004711	0. 77 0. 73
	HEMBA1004711	F-HEMBA1004715	0. 73
	HEMBA1004734	F-HEMBA1004734	0. 94
	HEMBA1004734	F-HEMBA1004734	0. 94
cc	HEMBA1004751	F-HEMBA1004751	0. 64
55	HEMBA1004752	F-HEMBA1004752	0. 87
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	HEMBA1004753	F-HEMBA1004753 ·	0. 54
	HEMBA1004756	F-HEMBA1004756	0. 54
_	HEMBA1004758	F-HEMBA1004758	0.69
5	HEMBA1004763	F-HEMBA1004763	0. 94
	HEMBA1004768	F-HEMBA1004768	0.66
	HEMBA1004771	F-HEMBA1004771	0.72
	HEMBA1004776	F-HEMBA1004776	0.49
10	HEMBA1004795	F-HEMBA1004795	0. 46
	HEMBA1004806	F-HEMBA1004806	0. 32
	HEMBA1004847	F-HEMBA1004847	0. 79
	HEMBA1004850	F-HEMBA1004850	0. 94
15	HEMBA1004863	F-HEMBA1004863	0. 32
	HEMBA1004864	F-HEMBA1004864	0.94
	HEMBA1004889	F-HEMBA1004889,	0. 94
	HEMBA1004923	F-HEMBA1004923	0.70
20	HEMBA1004929	F-HEMBA1004929	0.31
	HEMBA1004930	F-HEMBA1004930	0.90
	HEMBA1004933	F-HEMBA1004933	0.84
	HEMBA1004934	F-HEMBA1004934	0.34
25	HEMBA1004944	F-HEMBA1004944	0.40
	HEMBA1004954	F-HEMBA1004954	0. 62
	HEMBA1004972	F-HEMBA1004972	0. 90
	HEMBA1004973	F-HEMBA1004973	0.94
30	HEMBA1004977	F-HEMBA1004977	0.77
	HEMBA1004980	F-HEMBA1004980	0. 33
	HEMBA1005009	F-HEMBA1005009	0.94
	HEMBA1005019	F-HEMBA1005019	0. 32
35	HEMBA1005029	F-HEMBA1005029	0. 84
33	HEMBA1005035	F-HEMBA1005035	0. 91
	HEMBA1005047	F-HEMBA1005047	0. 94
	HEMBA1005050	F-HEMBA1005050	0. 88
	HEMBA1005066	F-HEMBA1005066	0. 79
40	HEMBA1005075	F-HEMBA1005075	0. 31
	HEMBA1005079	F-HEMBA1005079	0. 35
	HEMBA1005083	F-HEMBA1005083	0. 43
	HEMBA1005101	F-HEMBA1005101	0. 94
45	HEMBA1005113	F-HEMBA1005113	0. 38
	HEMBA1005133	F-HEMBA1005133	0.34
	HEMBA1005149	F-HEMBA1005149	0. 36
	HEMBA1005185	F-HEMBA1005185	0. 64
50	HEMBA1005201	F-HEMBA1005201	0. 83
	HEMBA1005202	F-HEMBA1005202	0. 94
	HEMBA1005206	F-HEMBA1005206	0. 94
	HEMBA1005219	F-HEMBA1005219	0. 57
55	HEMBA1005223	F-HEMBA1005223	0. 33
	HEMBA1005244	F-HEMBA1005244	0. 35

	HEMBA1005252	F-HEMBA1005252	0. 80
	HEMBA1005296	F-HEMBA1005296	0. 94
	HEMBA1005314	F-HEMBA1005314	0. 94
5	HEMBA1005331	F-HEMBA1005331	
	HEMBA1005331	F-HEMBA1005331	0. 94
	HEMBA1005367	F-HEMBA1005367	0. 50
			0. 91
10	HEMBA1005382	F-HEMBA1005382	0. 45
-	HEMBA1005394	F-HEMBA1005394	
	HEMBA1005403	F-HEMBA1005403	0. 94
	HEMBA1005411	F-HEMBA1005411	0. 33
45	HEMBA1005423	F-HEMBA1005423	0. 64
15	HEMBA1005468	F-HEMBA1005468	0. 33
	HEMBA1005469		0. 59
	HEMBA1005472		
		F-HEMBA1005474	0.51
20	HEMBA1005475	F-HEMBA1005475	0.36
	HEMBA1005513	F-HEMBA1005513	0.94
	HEMBA1005517	F-HEMBA1005517	0. 94
	HEMBA1005518	F-HEMBA1005518	0.39
25	HEMBA1005526	F-HEMBA1005526	0. 62
	HEMBA1005528	F-HEMBA1005528	0.81
	HEMBA1005530	F-HEMBA1005530	0. 62
	HEMBA1005548	F-HEMBA1005548	0.94
30	HEMBA1005558	F-HEMBA1005558	0. 94
	HEMBA1005576	F-HEMBA1005576	0. 49
	HEMBA1005581	F-HEMBA1005581	0. 53
	HEMBA1005582	F-HEMBA1005582	0. 34
35	HEMBA1005583	F-HEMBA1005583	0.94
	HEMBA1005595	F-HEMBA1005595	0. 93
	HEMBA1005609	F-HEMBA1005609	0.34
	HEMBA1005621	F-HEMBA1005621	0. 94
40	HEMBA1005666	F-HEMBA1005666	0. 57
40	HEMBA1005680	F-HEMBA1005680	0. 38
	HEMBA1005685	F-HEMBA1005685	0. 32
	HEMBA1005732	F-HEMBA1005732	0. 33
		F-HEMBA1005737	0. 92
45	HEMBA1005746		0. 37
	HEMBA1005755	F-HEMBA1005755	0. 94
	HEMBA1005813	F-HEMBA1005813	0. 33
	HEMBA1005815	F-HEMBA1005815	0. 82
50	HEMBA1005822	F-HEMBA1005822	0. 83
	HEMBA1005834	F-HEMBA1005834	0. 70
	HEMBA1005852	F-HEMBA1005852	0.76
	HEMBA1005884	F-HEMBA1005884	0. 79
55	HEMBA1005891	F-HEMBA1005891	0. 94
	HEMBA1005963	F-HEMBA1005963	0. 40

	HEMBA1005990	F-HEMBA1005990	0. 82
	HEMBA1005991	F-HEMBA1005991	0. 43
	HEMBA1006031	F-HEMBA1006031	0. 55
5	HEMBA1006035	F-HEMBA1006035	
	HEMBA1006081	F-HEMBA1006081	0. 36
	HEMBA1006091	F-HEMBA1006091	0. 81
	HEMBA1006100	F-HEMBA1006100	0. 31
10		F-HEMBA1006108	0. 80
	HEMBA1006108	F-HEMBA1006121	0. 45
	HEMBA1006121	F-HEMBA1006121	0. 45
	HEMBA1006130	F-HEMBA1006138	0. 48
15	HEMBA1006138		0. 48
15	HEMBA1006155	F-HEMBA1006155	0. 72
	HEMBA1006173	F-HEMBA1006173	
	HEMBA1006182	F-HEMBA1006182,	
	HEMBA1006198	F-HEMBA1006198 F-HEMBA1006235	0. 94 0. 58
20	HEMBA1006235		
	HEMBA1006248	F-HEMBA1006248 F-HEMBA1006252	0. 57 0. 31
	HEMBA1006252 HEMBA1006272	F-HEMBA1006272	0. 54
	HEMBA1006272	F-HEMBA1006272	0. 92
25	HEMBA1006283	F-HEMBA1006283	0. 90
	HEMBA1006284	F-HEMBA1006284	0. 93
	HEMBA1006291	F-HEMBA1006291	0. 38
	HEMBA1006293	F-HEMBA1006293	0. 89
30	HEMBA1006309	F-HEMBA1006309	0. 94
	HEMBA1006310	F-HEMBA1006310	0. 67
	HEMBA1006344	F-HEMBA1006344	0. 94
	HEMBA1006347	F-HEMBA1006347	0. 94
35	HEMBA1006349	F-HEMBA1006349	0. 56
	HEMBA1006377	F-HEMBA1006377	0. 94
	HEMBA1006381	F-HEMBA1006381	0. 33
	HEMBA1006398	F-HEMBA1006398	0.71
40	HEMBA1006424	F-HEMBA1006424	0. 40
	HEMBA1006445	F-HEMBA1006445	0.86
	HEMBA1006467	F-HEMBA1006467	0. 55
	HEMBA1006474	F-HEMBA1006474	0.66
45	HEMBA1006483	F-HEMBA1006483	0. 71
	HEMBA1006485	F-HEMBA1006485	0.87
	HEMBA1006492	F-HEMBA1006492	0. 73
	HEMBA1006494	F-HEMBA1006494	0.90
50	HEMBA1006497	F-HEMBA1006497	0.77
•	HEMBA1006502	F-HEMBA1006502	0.80
	HEMBA1006507	F-HEMBA1006507	0.56
	HEMBA1006521	F-HEMBA1006521	0. 91
55	HEMBA1006530	F-HEMBA1006530	0.37
55	HEMBA1006535	F-HEMBA1006535	056

	HEMBA1006559	F-HEMBA1006559	0.39
	HEMBA1006566	F-HEMBA1006566	0. 41
	HEMBA1006579	F-HEMBA1006579	0.83
5	HEMBA1006583	F-HEMBA1006583	0.31
	HEMBA1006612	F-HEMBA1006612	0. 55
	HEMBA1006624	F-HEMBA1006624	0.94
	HEMBA1006631	F-HEMBA1006631	0. 74
10	HEMBA1006643	F-HEMBA1006643	0. 38
	HEMBA1006652	F-HEMBA1006652	0. 93
	HEMBA1006659	F-HEMBA1006659	0.63
	HEMBA1006674	F-HEMBA1006674	0.54
15	HEMBA1006682	F-HEMBA1006682	0.34
	HEMBA1006708	F-HEMBA1006708	0.40
	HEMBA1006709	F-HEMBA1006709,	0. 41
	HEMBA1006717	F-HEMBA1006717	0. 43
20	HEMBA1006737	F-HEMBA1006737	0.72
	HEMBA1006754	F-HEMBA1006754	·0. 45
	HEMBA1006758	F-HEMBA1006758	0.60
	HEMBA1006767	F-HEMBA1006767	0.61
25	HEMBA1006789	F-HEMBA1006789	0.34
	HEMBA1006795	F-HEMBA1006795	0.48
	HEMBA1006796	F-HEMBA1006796	0.41
	HEMBA1006807	F-HEMBA1006807	0.34
30	HEMBA1006832	F-HEMBA1006832	0.59
30	HEMBA1006877	F-HEMBA1006877	0.90
	HEMBA1006885	F-HEMBA1006885	0. 80
	HEMBA1006900	F-HEMBA1006900	0.34
	HEMBA1006914	F-HEMBA1006914	0. 78
35	HEMBA1006926	F-HEMBA1006926	0.67
	HEMBA1006936	F-HEMBA1006936	0. 65
	HEMBA1006941	F-HEMBA1006941	0. 94
	HEMBA1006973	F-HEMBA1006973	0. 64
40	HEMBA1006976	F-HEMBA1006976	0.40
	HEMBA1006993	F-HEMBA1006993	0.81
	HEMBA1006996	F-HEMBA1006996	0. 77
	HEMBA1007002	F-HEMBA1007002	0. 72
45	HEMBA1007018	F-HEMBA1007018	0. 31
	HEMBA1007052	F-HEMBA1007052	0. 41
	HEMBA1007062	F-HEMBA1007062	0.39
	HEMBA1007066	F-HEMBA1007066	0. 35
50	HEMBA1007080	F-HEMBA1007080	0. 43
	HEMBA1007085	F-HEMBA1007085	0. 73
	HEMBA1007087	F-HEMBA1007087	0. 94
	HEMBA1007112	F-HEMBA1007112	0.49
55	HEMBA1007121	F-HEMBA1007121	0.70
	HEMBA1007149	F-HEMBA1007149	0. 94

	HEMBA1007151	F-HEMBA1007151	0. 80
	HEMBA1007174	F-HEMBA1007174	0.63
	HEMBA1007178	F-HEMBA1007178	0. 47
5	HEMBA1007194	F-HEMBA1007194	0.94
	HEMBA1007203	F-HEMBA1007203	0. 48
	HEMBA1007224	F-HEMBA1007224	0.74
	HEMBA1007243	F-HEMBA1007243	0. 94
10	HEMBA1007251	F-HEMBA1007251	0. 79
	HEMBA1007267	F-HEMBA1007267	0.86
	HEMBA1007281	F-HEMBA1007281	0.61
	HEMBA1007300	F-HEMBA1007300	0. 87
15	HEMBA1007301	F-HEMBA1007301	0. 38
	HEMBA1007319	F-HEMBA1007319	0. 47
	HEMBA1007320	F-HEMBA1007320,	0.69
	HEMBA1007342	F-HEMBA1007342	0.80
20	HEMBB1000008	F-HEMB81000008	0.39
	HEMBB1000018	F-HEMBB1000018	0. 79
	HEMBB1000024	F-HEMBB1000024	0.62
	HEMBB1000025	F-HEMBB1000025	0.66
25	HEMBB1000030	F-HEMBB1000030	0. 33
	HEMBB1000036	F-HEMBB1000036	0. 75
	HEMBB1000037	F-HEMBB1000037	0. 94
	HEMBB1000048	F-HEMBB1000048	0. 87
30	HEMBB1000083	F-HEMBB1000083	0. 48
	HEMBB1000103	F-HEMBB1000103	0. 63
	HEMBB1000119	F-HEMBB1000119	0. 94
	HEMBB1000136	F-HEMBB1000136	0. 92
35	HEMBB1000173	F-HEMBB1000173	0. 46
•	HEMBB1000175	F-HEMBB1000175	0. 48
	HEMBB1000198	F-HEMBB1000198	0. 74
	HEMBB1000215	F-HEMBB1000215	0.61
40	HEMBB1000217 HEMBB1000226	F-HEMBB1000217 F-HEMBB1000226	0. 94 0. 94
70	HEMBB1000240	F-HEMBB1000240	0. 40
	HEMBB1000240	F-HEMBB1000244	0. 40
	HEMBB1000244	F-HEMBB1000266	0. 92
45	HEMBB1000338	F-HEMBB1000338	0. 33
45	HEMBB1000339	F-HEMBB1000339	0. 40
	HEMBB1000391	F-HEMBB1000391	0. 43
	HEMBB1000438	F-HEMBB1000438	0. 79
50	HEMBB1000449	F-HEMBB1000449	0. 48
50	HEMBB1000510	F-HEMBB1000510	0. 35
	HEMBB1000550	F-HEMBB1000550	0. 94
	HEMBB1000556	F-HEMBB1000556	0. 89
	HEMBB1000589	F-HEMBB1000589	0. 77
55	HEMBB1000591	F-HEMBB1000591	0. 33

	HEMBB1000592	F-HEMBB1000592	0.77
	HEMBB1000593	F-HEMBB1000593	0.68
	HEMBB1000623	F-HEMBB1000623	0.94
5	HEMBB1000630	F-HEMBB1000630	0.59
	HEMBB1000631	F-HEMBB1000631	0.71
	HEMBB1000632	F-HEMBB1000632	0.38
	HEMBB1000671	F-HEMBB1000671	0.36
10	HEMBB1000673	F-HEMBB1000673	0.54
	HEMBB1000693	F-HEMBB1000693	0.68
	HEMBB1000705	F-HEMBB1000705	0.44
	HEMBB1000706	F-HEMBB1000706	0.45
15	HEMBB1000725	F-HEMBB1000725	0.91
	HEMBB1000763	F-HEMBB1000763	0.87
	HEMBB1000774	F-HEMBB1000774	0.80
	HEMBB1000781	F-HEMBB1000781	0.94
20	HEMBB1000789	F-HEMBB1000789	0.34
	HEMBB1000807	F-HEMBB1000807	0.94
	HEMBB1000810	F-HEMBB1000810	0.33
	HEMBB1000826	F-HEMBB1000826	0.36
25	HEMBB1000835	F-HEMBB1000835	0. 81
	HEMBB1000848	F-HEMBB1000848	0.66
	HEMBB1000852	F-HEMBB1000852	0.85
	HEMBB1000870	F-HEMB81000870	0.34
30	HEMBB1000887	F-HEMBB1000887	0.58
30	HEMBB1000908	F-HEMBB1000908	0.31
	HEMBB1000927	F-HEMBB1000927	0.94
	HEMBB1000947	F-HEMBB1000947	0. 33
	HEMBB1000973	F-HEMBB1000973	0.72
35	HEMBB1000975	F-HEMBB1000975	0.68
	HEMBB1000985	F-HEMBB1000985	0.62
	HEMBB1000991	F-HEMBB1000991	0.45
	HEMBB1001011	F-HEMBB1001011	0. 58 0. 86
40	HEMBB1001014	F-HEMBB1001014	
	HEMBB1001020	F-HEMBB1001020	0.87
	HEMBB1001024 HEMBB1001056	F-HEMBB1001024 F-HEMBB1001056	0. 64 0. 94
	HEMBB1001058	F-HEMBB1001058	0. 39
45	HEMBB1001068	F-HEMBB1001068	0.57
	HEMBB1001008	F-HEMBB1001096	0. 48
	HEMB8100105	F-HEMBB1001105	0. 45
	HEMBB1001112	F-HEMBB1001112	0. 47
50	HEMBB1001117	F-HEMBB1001117	0.83
	HEMBB1001126	F-HEMBB1001126	0.76
	HEMBB1001137	F-HEMBB1001137	0. 53
	HEMBB1001151	F-HEMBB1001151	0.94
55	HEMBB1001153	F-HEMBB1001153	0.77

	HEMBB1001169 F-HEMBB1001169 0	. 39
	HEMBB1001175 F-HEMBB1001175 0	). 91
	HEMBB1001182 F~HEMBB1001182 0	. 94
5		. 54
		. 88
		. 67
		. 33
10		. 33
		. 58
		. 86
		. 76
15	HEMBB1001331 F-HEMBB1001331 0	. 44
		. 40
	HEMBB1001346 F-HEMBB1001346, 0	. 83
	No. of the contract of the con	. 69
20	HEMBB1001369 F-HEMBB1001369 0	. 47
	HEMBB1001384 - F-HEMBB1001384 0	. 80
	HEMBB1001387 F-HEMBB1001387 0	. 36
	HEMBB1001394 F-HEMBB1001394 0	. 94
25	HEMBB1001410 F-HEMBB1001410 0	. 51
	HEMBB1001426 F-HEMBB1001426 0	. 80
	HEMBB1001429 F-HEMBB1001429 0	. 91
		. 31
30		. 44
		. 58
		. 65
		. 46
35		. 74
33		. 81
		. 46
		. 33
40		. 39
40		. 38
		. 83
	HEMBB1001684 F-HEMBB1001684 0	
		. 90
45		. 92
		. 51
		. 94
		. 88 . 74
50		. 33
		. 41
		. 93
		. 78
55		. 32
	######################################	. 52

	HEMBB1001869	F-HEMBB1001869	0. 48
	HEMBB1001872	F-HEMBB1001872	0. 93
	HEMBB1001874	F-HEMBB1001874	0.31
5	HEMBB1001905	F-HEMBB1001905	0. 87
	HEMBB1001906	F-HEMBB1001906	0. 88
	HEMBB1001908	F-HEMBB1001908	0. 69
	HEMBB1001910	F-HEMBB1001910	0. 77
10	HEMBB1001915	F-HEMBB1001915	0. 76
	HEMBB1001945	F-HEMBB1001945	0. 64
	HEMBB1001947	F-HEMBB1001947	0. 94
	HEMBB1001950	F-HEMBB1001950	0. 73
15	HEMBB1001957	F-HEMBB1001957	0. 64
	HEMBB1001962	F-HEMBB1001962	0. 90
	HEMBB1001990	F-HEMBB1001990,	
	HEMBB1002044	F-HEMBB1002044	0. 82
20	HEMBB1002050	F-HEMBB1002050	0. 91
	HEMBB1002068	F-HEMBB1002068	0. 48
	HEMBB1002134	F-HEMBB1002134	0.80
	HEMBB1002142	F-HEMBB1002142	0.39
26	HEMBB1002152	F-HEMBB1002152	0.31
25	HEMBB1002193	F-HEMBB1002193	0. 35
	HEMBB1002217	F-HEMBB1002217	0.49
	HEMBB1002218	F-HEMBB1002218	0. 32
	HEMBB1002249	F-HEMBB1002249	0.72
30	HEMBB1002300	F-HEMBB1002300	0. 75
	HEMBB1002327	F-HEMBB1002327	0.86
	HEMBB1002329	F-HEMBB1002329	0.31
	HEMBB1002340	F-HEMBB1002340	0. 42
35	HEMBB1002342	F-HEMBB1002342	0. 76
	HEMBB1002358	F-HEMBB1002358	0.70
	HEMBB1002359	F-HEMBB1002359	0. 75
	HEMBB1002383	F-HEMBB1002383	0. 78
40	HEMBB1002409	F-HEMBB1002409	0. 52
	HEMBB1002415	F-HEMBB1002415	0.82
	HEMBB1002442	F-HEMB81002442	0. 31
	HEMBB1002457	F-HEMBB1002457	0. 46
45	HEMBB1002489	F-HEMBB1002489	0. 69
	HEMBB1002492	F-HEMBB1002492	0. 68
	HEMBB1002495	F-HEMBB1002495	0. 94
	HEMBB1002502	F-HEMBB1002502	0.39
50		F-HEMBB1002510	0. 94
	HEMBB1002545	F-HEMBB1002545	0. 44
	HEMBB1002550	F-HEMB81002550	0. 57
	HEMBB1002600	F-HEMBB1002600	0.91
55	HEMBB1002607	F-HEMB81002607	0. 83
55	HEMBB1002614	F-HEMBB1002614	0. 75

	HEMBB1002684	F-HEMBB1002684	0.84
	HEMBB1002692	F-HEMBB1002692	0. 52
	HEMBB1002697	F-HEMBB1002697	0.86
5	HEMBB1002705	F-HEMBB1002705	0.32
	MAMMA1000019	F-MAMMA1000019	0. 48
	MAMMA1000020	F-MAMMA1000020	0. 36
	MAMMA1000025	F-MAMMA1000025	0. 31
10	MAMMA1000055	F-MAMMA1000055	0.39
	MAMMA1000057	F-MAMMA1000057	0.35
	MAMMA1000069	F-MAMMA1000069	0. 78
	MAMMA1000084	F-MAMMA1000084	0. 61
15	MAMMA1000085	F-MAMMA1000085	0.72
	MAMMA1000139	F-MAMMA1000139	0.33
	MAMMA1000143	F-MAMMA1000143	
	MAMMA1000163	F-MAMMA1000163	0.94
20	MAMMA1000171	F-MAMMA1000171	0. 78
	MAMMA1000173		
	MAMMA1000183		0. 77
	MAMMA1000251		0. 94
25	MAMMA1000257		0. 75
	MAMMA1000277	F-MAMMA1000277	0. 94
	MAMMA1000278	F-MAMMA1000278	0. 35
	MAMMA1000279 MAMMA1000284	F-MAMMA1000279	0. 62
30		F-MAMMA1000284	0.35
	MAMMA1000309 MAMMA1000312	F-MAMMA1000309	0.53
	MAMMA1000312	F-MAMMA1000312 F-MAMMA1000313	0.68
	MAMMA1000313	F-MAMMA1000313	0. 32 0. 35
35	MAMMA1000359	F-MAMMA1000361	0. 61
	MAMMA1000372	F-MAMMA1000372	0.60
	MAMMA1000388	F-MAMMA1000388	0. 93
	MAMMA1000395	F-MAMMA1000395	0. 94
40	MAMMA1000410	F-MAMMA1000410	0. 94
	MAMMA1000416	F-MAMMA1000416	0.89
	MAMMA1000421	F-MAMMA1000421	0. 50
	MAMMA1000422	F-MAMMA1000422	0.92
45	MAMMA1000429	F-MAMMA1000429	0.92
	MAMMA1000458	F-MAMMA1000458	0.88
	MAMMA1000468	F-MAMMA1000468	0.60
	MAMMA1000472	F-MAMMA1000472	0. 33
50	MAMMA1000490	F-MAMMA1000490	0.34
	MAMMA1000524	F-MAMMA1000524	0.39
	MAMMA1000567	F-MAMMA1000567	0. 40
	MAMMA1000583	F-MAMMA1000583	0.45
	MAMMA1000612	F-MAMMA1000612	0.89
55	MAMMA1000623	F-MAMMA1000623	0. 44
		•	

		MAMMA1000625	F-MAMMA1000625	0. 52
		MAMMA1000664	F-MAMMA1000664	0. 42
		MAMMA1000670	F-MAMMA1000670	0. 94
	5	MAMMA1000672	F-MAMMA1000672	0.44
		MAMMA1000684	F-MAMMA1000684	0.94
		MAMMA1000713	F-MAMMA1000713	0. 88
		MAMMA1000731	F-MAMMA1000731	0. 67
;	10	MAMMA1000734	F-MAMMA1000734	0.40
		MAMMA1000738	F-MAMMA1000738	0. 93
		MAMMA1000746	F-MAMMA1000746	0.48
		MAMMA1000760	F-MAMMA1000760	0.94
1	15	MAMMA1000776	F-MAMMA1000776	0.94
		MAMMA1000824	F-MAMMA1000824	0. 81
		MAMMA1000831	F-MAMMA1000831	0. 39
		MAMMA1000841	F-MAMMA1000841	0.41
2	20	MAMMA1000842	F-MAMMA1000842	0.61
		MAMMA1000843	F-MAMMA1000843	0.48
		MAMMA1000856	F-MAMMA1000856	0.31
		MAMMA1000859	F-MAMMA1000859	0. 44
2	25	MAMMA1000865	F-MAMMA1000865	0. 94
		MAMMA1000875	F-MAMMA1000875	0. 49
		MAMMA1000897	F-MAMMA1000897	0. 94
		MAMMA1000906	F-MAMMA1000906	0. 45
3	0	MAMMA1000908	F-MAMMA1000908	0. 39
		MAMMA1000914	F-MAMMA1000914	0. 94
		MAMMA1000921	F-MAMMA1000921	0. 34
		MAMMA1000956	F-MAMMA1000956	0. 39
3	5	MAMMA1000968	F-MAMMA1000968	0. 35
		MAMMA1000979 MAMMA1001008	F-MAMMA1000979 F-MAMMA1001008	0. 68
		MAMMA1001008	F-MAMMA1001003	0. 36 0. 48
		MAMMA1001021	F-MAMMA1001041	0. 46
4	0	MAMMA1001059	F-MAMMA1001059	0. 94
		MAMMA1001033	F-MAMMA1001073	0. 94
		MAMMA1001075	F-MAMMA1001075	0. 94
		MAMMA1001078	F-MAMMA1001078	0. 52
4	5	MAMMA1001080	F-MAMMA1001080	0. 87
•		MAMMA1001091	F-MAMMA1001091	0. 53
		MAMMA1001105	F-MAMMA1001105	0. 94
		MAMMA1001110	F-MAMMA1001110	0. 78
5	0	MAMMA1001126	F-MAMMA1001126	0. 85
3	-	MAMMA1001139	F-MAMMA1001139	0. 38
		MAMMA1001143	F-MAMMA1001143	0. 79
		MAMMA1001154	F-MAMMA1001154	0. 31
,,	=	MAMMA1001181	F-MAMMA1001181	0.69
5	,	MAMMA1001198	F-MAMMA1001198	0.88

	MAMMA1001202	F-MAMMA1001202	0.38
	MAMMA1001215	F-MAMMA1001215	0.94
	MAMMA1001222	F-MAMMA1001222	0.89
5	MAMMA1001244	F-MAMMA1001244	0.87
	MAMMA1001259	F-MAMMA1001259	0. 91
	MAMMA1001260	F-MAMMA1001260	0. 75
	MAMMA1001271	F-MAMMA1001271	0.77
10	MAMMA1001292	F-MAMMA1001292	0.94
	MAMMA1001296	F-MAMMA1001296	0.75
	MAMMA1001305	F-MAMMA1001305	0.66
	MAMMA1001343	F-MAMMA1001343	0.38
15	MAMMA1001346	F-MAMMA1001346	0.81
	MAMMA1001388	F-MAMMA1001388	0. 94
	MAMMA1001411	F-MAMMA1001411,	
	MAMMA1001411	F-MAMMA1001411	0. 42
20	MAMMA1001465	F-MAMMA1001465	0. 94
	MAMMA1001476	F-MAMMA1001476	0. 90
	MAMMA1001487	F-MAMMA1001487	0. 78
	MAMMA1001510	F-MAMMA1001510	0. 78
25	MAMMA1001522	F-MAMMA1001522	0. 63
25	MAMMA1001551	F-MAMMA1001551	0. 81
	MAMMA1001576		0. 94
	MAMMA1001600	F-MAMMA1001600	0. 72
	MAMMA1001604	F-MAMMA1001604	0. 72
30	MAMMA1001620	F-MAMMA1001620	0. 43
	MAMMA1001627	F-MAMMA1001627	0. 39
	MAMMA1001630	F-MAMMA1001630	0. 46
	MAMMA1001633	F-MAMMA1001633	0. 58
35	MAMMA1001635	F-MAMMA1001635	0. 89
	MAMMA1001649	F-MAMMA1001649	0.62
	MAMMA1001654	F-MAMMA1001654	0. 94
	MAMMA1001692	F-MAMMA1001692	0. 83
40	MAMMA1001730	F-MAMMA1001730	0. 48
	MAMMA1001735	F-MAMMA1001735	0. 94
	MAMMA1001743	F-MAMMA1001743	0. 94
	MAMMA1001744	F-MAMMA1001744	0. 33
45	MAMMA1001751	F-MAMMA1001751	0. 94
	MAMMA1001754	F-MAMMA1001754	0. 47
	MAMMA1001757	F-MAMMA1001757	0. 38
	MAMMA1001764	F-MAMMA1001764	0. 57
	MAMMA1001768	F-MAMMA1001768	0. 85
50	MAMMA1001771	F-MAMMA1001771	0. 39
	MAMMA1001771	F-MAMMA1001771	
	MAMMA1001785	F-MAMMA1001785	0.61
	MAMMA1001789	F-MAMMA1001790	0.41
55	MAMMA1001790	F-MAMMA1001812	0.69
	III/WIII/N I UU I O I Z	I WARMA LOUIO (Z	0.32

	MAMMA1001817	F-MAMMA1001817	0. 55
	MAMMA1001824	F-MAMMA1001824	0.57
	MAMMA1001837	F-MAMMA1001837	0. 35
5	MAMMA1001848	F-MAMMA1001848	
	MAMMA1001851	F-MAMMA1001851	0, 46
	MAMMA1001858	F-MAMMA1001858	0. 53
	MAMMA1001864	F-MAMMA1001864	0. 32
10	MAMMA1001868	F-MAMMA1001868	0.48
	MAMMA1001874	F-MAMMA1001874	0. 94
	MAMMA1001956	F-MAMMA1001956	0.47
	MAMMA1001969	F-MAMMA1001969	0, 31
15	MAMMA1001970	F-MAMMA1001970	0.34
	MAMMA1002009	F-MAMMA1002009	0.45
	MAMMA1002011	F-MAMMA1002011,	
	MAMMA1002033	F-MAMMA1002033	0.33
20	MAMMA1002041	F-MAMMA1002041	0.86
	MAMMA1002042	F-MAMMA1002042	0.88
	MAMMA1002047	F-MAMMA1002047	0. 41
	MAMMA1002068	F-MAMMA1002068	0. 38
25	MAMMA1002143	F-MAMMA1002143	0. 91
20	MAMMA1002153	F-MAMMA1002153	0.94
	MAMMA1002156	F-MAMMA1002156	0.73
	MAMMA1002170	F-MAMMA1002170	0.43
	MAMMA1002174	F-MAMMA1002174	0. 69
30	MAMMA1002198	F-MAMMA1002198	0.65
	MAMMA1002209	F-MAMMA1002209	0.80
	MAMMA1002219	F-MAMMA1002219	0. 85
	MAMMA1002236	F-MAMMA1002236	0. 55
35	MAMMA1002243	F-MAMMA1002243	0. 50
	MAMMA1002268	F-MAMMA1002268	0. 43
	MAMMA1002269	F-MAMMA1002269	0. 33
	MAMMA1002292	F-MAMMA1002292	0. 76
40	MAMMA1002294	F-MAMMA1002294	0.94
	MAMMA1002297	F-MAMMA1002297	0. 91
	MAMMA1002308.	F-MAMMA1002308	0. 68
	MAMMA1002312	F-MAMMA1002312	0.77
45	MAMMA1002317	F-MAMMA1002317	0. 46
	MAMMA1002319	F-MAMMA1002319	0. 31
	MAMMA1002329	F-MAMMA1002329	0. 34
	MAMMA1002333	F-MAMMA1002333	0. 65
50	MAMMA1002351	F-MAMMA1002351	0.89
	MAMMA1002353	F-MAMMA1002353	0. 94
	MAMMA1002355	F-MAMMA1002355	0. 55
	MAMMA1002356	F-MAMMA1002356	0. 32
55	MAMMA1002362	F-MAMMA1002362	0.83
55	MAMMA1002380	F-MAMMA1002380	0. 94

	MAMMA1002384	F-MAMMA1002384	0. 42
	MAMMA1002385	F-MAMMA1002385	0. 82
_	MAMMA1002413	F-MAMMA1002413	0.94
5	MAMMA1002427	F-MAMMA1002427	0.40
	MAMMA1002434	F-MAMMA1002434	0. 74
	MAMMA1002454	F-MAMMA1002454	0.82
	MAMMA1002461	F-MAMMA1002461	0.49
10	MAMMA1002470	F-MAMMA1002470	0.89
	MAMMA1002485	F-MAMMA1002485	0.55
	MAMMA1002524	F-MAMMA1002524	0.33
	MAMMA1002530	F-MAMMA1002530	0.75
15	MAMMA1002554	F-MAMMA1002554	0.41
	MAMMA1002585	F-MAMMA1002585	0. 31
	MAMMA1002590	F-MAMMA1002590,	0.72
	MAMMA1002598	F-MAMMA1002598	0. 76
20	MAMMA1002617	F-MAMMA1002617	0.49
	MAMMA1002619	F-MAMMA1002619	0. 88
	MAMMA1002637	F-MAMMA1002637	0. 94
	MAMMA1002650	F-MAMMA1002650	0. 54
25	MAMMA1002655	F-MAMMA1002655	0. 39
	MAMMA1002665	F-MAMMA1002665	0. 33
	MAMMA1002671	F-MAMMA1002671	0. 94
	MAMMA1002673	F-MAMMA1002673	0.70
30	MAMMA1002684	F-MAMMA1002684	0. 94
	MAMMA1002685	F-MAMMA1002685	0. 88
	MAMMA1002699	F-MAMMA1002699	0.94
	MAMMA1002711 MAMMA1002769	F-MAMMA1002711 F-MAMMA1002769	0. 39 0. 88
35	MAMMA1002775		0. 81
	MAMMA1002782	F-MAMMA10027782	0.87
	MAMMA1002796	F-MAMMA1002796	0. 40
	MAMMA1002807	F-MAMMA1002807	0. 64
40	MAMMA1002842	F-MAMMA1002842	0. 94
	MAMMA1002843	F-MAMMA1002843	0. 32
	MAMMA1002868	F-MAMMA1002868	0. 54
	MAMMA1002869	F-MAMMA1002869	0. 94
45	MAMMA1002881	F-MAMMA1002881	0. 94
	MAMMA1002886	F-MAMMA1002886	0. 61
	MAMMA1002890	F-MAMMA1002890	0. 94
	MAMMA1002895	F-MAMMA1002895	0.94
50	MAMMA1002937	F-MAMMA1002937	0. 91
	MAMMA1002938	F-MAMMA1002938	0. 78
	MAMMA1002947	F-MAMMA1002947	0.94
	MAMMA1002964	F-MAMMA1002964	0. 34
55	MAMMA1002987	F-MAMMA1002987	0.74
	MAMMA1003011	F-MAMMA1003011	0.90

		MAMMA1003013	F-MAMMA1003013	0.36
		MAMMA1003015	F-MAMMA1003015	0. 92
		MAMMA1003035	F-MAMMA1003035	0.94
	5	MAMMA1003044	F-MAMMA1003044	
		MAMMA1003047	F-MAMMA1003047	0. 94
		MAMMA1003049	F-MAMMA1003049	0. 31
		MAMMA1003056	F-MAMMA1003056	0. 75
	10	MAMMA1003057	F-MAMMA1003057	0. 33
		MAMMA1003066	F-MAMMA1003066	0.38
		MAMMA1003099	F-MAMMA1003099	0. 79
		MAMMA1003104	F-MAMMA1003104	0.44
	15	MAMMA1003113	F-MAMMA1003113	0. 90
		MAMMA1003127	F-MAMMA1003127	0. 53
		MAMMA1003135	F-MAMMA1003135,	0. 94
		MAMMA1003146	F-MAMMA1003146	0. 88
	20	MAMMA1003150	F-MAMMA1003150	0.73
		MAMMA1003166	F-MAMMA1003166	0. 94
		NT2RM1000001	F-NT2RM1000001	0. 66
		NT2RM1000018	F-NT2RM1000018	0. 55
	25	NT2RM1000032	F-NT2RM1000032	0. 74
		NT2RM1000035	F-NT2RM1000035	0. 52
		NT2RM1000037	F-NT2RM1000037	0. 94
		NT2RM1000039	F-NT2RM1000039	0. 77
	30	NT2RM1000055	F-NT2RM1000055	0. 89
		NT2RM1000059	F-NT2RM1000059	0. 41
		NT2RM1000062	F-NT2RM1000062	0. 65
		NT2RM1000080	F-NT2RM1000080	0. 61
	35	NT2RM1000086	F-NT2RM1000086	0. 94
		NT2RM1000092	F-NT2RM1000092	0. 94
		NT2RM1000118	F-NT2RM1000118	0. 76
		NT2RM1000119	F-NT2RM1000119	0. 41
	40	NT2RM1000127	F-NT2RM1000127	0. 94
	<del>4</del> 0	NT2RM1000131	F-NT2RM1000131	0. 90
		NT2RM1000132	F-NT2RM1000132	0. 94
		N12RM1000153	F-NT2RM1000153	0. 90
		NT2RM1000186	F-NT2RM1000186	0. 78
	45	NT2RM1000187	F-NT2RM1000187	0. 54
		NT2RM1000199	F-NT2RM1000199	0. 33
		NT2RM1000242	F-NT2RM1000242	0. 78
		NT2RM1000244	F-NT2RM1000244	0. 31
•	50	NT2RM1000252	F-NT2RM1000252	0. 76
		NT2RM1000256	F-NT2RM1000256	0.94
		NT2RM1000257	F-NT2RM1000257	0.89
		NT2RM1000260	F-NT2RM1000260	0.42
	55	NT2RM1000271 NT2RM1000272	F-NT2RM1000271	0. 74 0. 79
		111 2 Non 1 UUU 2 / 2	F-NT2RM1000272	0. 78

	NT2RM1000280	F-NT2RM1000280	0. 94
	NT2RM1000300	F-NT2RM1000300	0. 47
	NT2RM1000314	F-NT2RM1000314	0.61
5	NT2RM1000318	F-NT2RM1000318	0.94
	NT2RM1000341	F-NT2RM1000341	0. 70
	NT2RM1000354	F-NT2RM1000354	0. 82
	NT2RM1000355	F-NT2RM1000355	0. 94
10	NT2RM1000365	F-NT2RM1000365	0.31
	NT2RM1000377	F-NT2RM1000377	0.87
	NT2RM1000388	F-NT2RM1000388	0.94
	NT2RM1000394	F-NT2RM1000394	0. 58
15	NT2RM1000399	F-NT2RM1000399	0. 86
	NT2RM1000421	F-NT2RM1000421	0. 39
	NT2RM1000430		
	NT2RM1000499	F-NT2RM1000499	0.44
20	NT2RM1000539	F-NT2RM1000539	0.87
20	NT2RM1000553	F-NT2RM1000553	0.88
	NT2RM1000555	F-NT2RM1000555	0.37
	NT2RM1000563	F-NT2RM1000563	0. 63
05	NT2RM1000623	F-NT2RM1000623	0.34
25	NT2RM1000648	F-NT2RM1000648	0.94
	NT2RM1000661	F-NT2RM1000661	0.61
	NT2RM1000666	F-NT2RM1000666	0.94
	NT2RM1000669	F-NT2RM1000669	0.94
30	NT2RM1000672	F-NT2RM1000672	0.53
	NT2RM1000691	F-NT2RM1000691	0.38
	NT2RM1000699	F-NT2RM1000699	0.32
	NT2RM1000702	F-NT2RM1000702	0. 74
35	NT2RM1000725	F-NT2RM1000725	0. 75
	NT2RM1000741	F-NT2RM1000741	0. 94
	NT2RM1000742	F-NT2RM1000742	0. 54
	NT2RM1000746	F-NT2RM1000746	0.94
40	NT2RM1000770	F-NT2RM1000770	0. 48
	NT2RM1000772	F-NT2RM1000772	0.39
	NT2RM1000780	F-NT2RM1000780	0.66
	NT2RM1000781	F-NT2RM1000781	0.49
45	NT2RM1000800	F-NT2RM1000800	0. 72
	NT2RM1000802	F-NT2RM1000802	0.89
	NT2RM1000811	F-NT2RM1000811	0. 53
	NT2RM1000826	F-NT2RM1000826	0.39
50	NT2RM1000829	F-NT2RM1000829	0.63
	NT2RM1000833	F-NT2RM1000833	0.94
	NT2RM1000850	F-NT2RM1000850	0.89
	NT2RM1000852	F-NT2RM1000852	0.88
55	NT2RM1000857	F-NT2RM1000857	0.58
	NT2RM1000867	F-NT2RM1000867	0.90

	NT2RM1000874	F-NT2RM1000874	0.89
	NT2RM1000882	F-NT2RM1000882	0.63
	NT2RM1000883	F-NT2RM1000883	0.71
5	NT2RM1000885	F-NT2RM1000885	
	NT2RM1000894	F-NT2RM1000894	0. 57
	NT2RM1000898	F-NT2RM1000898	0. 87
	NT2RM1000905	F-NT2RM1000905	0.94
10	NT2RM1000924	F-NT2RM1000924	0. 88
	NT2RM1000927	F-NT2RM1000927	0.70
	NT2RM1000962	F-NT2RM1000962	0.84
	NT2RM1000978	F-NT2RM1000978	0. 39
15	NT2RM1001003	F-NT2RM1001003	0. 81
	NT2RM1001008	F-NT2RM1001008	0.94
	NT2RM1001043	F-NT2RM1001043,	
	NT2RM1001044	F-NT2RM1001044	0. 94
20	NT2RM1001059	F-NT2RM1001059	0.94
	NT2RM1001066	F-NT2RM1001066	0.94
	NT2RM1001072	F-NT2RM1001072	0.77
	NT2RM1001074	F-NT2RM1001074	0.94
25	NT2RM1001082	F-NT2RM1001082	0.94
	NT2RM1001085	F-NT2RM1001085	0.83
	NT2RM1001092	F-NT2RM1001092	0.77
	NT2RM1001102	F-NT2RM1001102	0.59
30	NT2RM1001105	F-NT2RM1001105	0.94
	NT2RM1001112	F-NT2RM1001112	0. 50
	NT2RM1001115	F-NT2RM1001115	0.32
	NT2RM1001139	F-NT2RM1001139	0.77
35	NT2RM2000006	F-NT2RM2000006	0.34
	NT2RM2000013	F-NT2RM2000013	0.70
	NT2RM2000030	F-NT2RM2000030	0.89
	NT2RM2000032	F-NT2RM2000032	0.36
40	NT2RM2000042	F-NT2RM2000042	0.32
,,	NT2RM2000092	F-NT2RM2000092	0.55
	NT2RM2000093 NT2RM2000101	F-NT2RM2000093	0.33
	NT2RM2000101	F-NT2RM2000101 F-NT2RM2000124	0. 77 0. 94
45	NT2RM2000124	F-NT2RM2000124	0. 94
43	NT2RM2000191	F-NT2RM2000191	0. 93
	NT2RM2000132	F-NT2RM2000132	0. 68
	NNNNNNNNNN	F-NNNNNNNNNNNN	0. 94
50	NT2RM2000250	F-NT2RM2000250	0. 94
50	NT2RM2000259	F-NT2RM2000259	0. 68
	NT2RM2000260	F-NT2RM2000260	0. 37
	NT2RM2000287	F-NT2RM2000287	0. 46
	NT2RM2000322	F-NT2RM2000322	0. 83
55	NT2RM2000359	F-NT2RM2000359	0. 94
			•

	NT2RM2000363	F-NT2RM2000363	0.91
	NT2RM2000368	F-NT2RM2000368	0.34
5	NT2RM2000371	F-NT2RM2000371	0.94
5	NT2RM2000374	F-NT2RM2000374_	0, 65
	NT2RM2000395	F-NT2RM2000395	0.33
	NT2RM2000402	F-NT2RM2000402	0.80
	NT2RM2000407	F-NT2RM2000407	0.88
10	NT2RM2000420	F-NT2RM2000420	0.92
	NT2RM2000422	F-NT2RM2000422	0.49
	NT2RM2000452	F-NT2RM2000452	0.93
	NT2RM2000469	F-NT2RM2000469	0.39
15	NT2RM2000490	F-NT2RM2000490	0.90
	NT2RM2000502	F-NT2RM2000502	0.64
	NT2RM2000504	F-NT2RM2000504	0.71
	NT2RM2000522	F-NT2RM2000522	0.91
20	NT2RM2000540	F-NT2RM2000540	0.87
	NT2RM2000556	F-NT2RM2000556	0.91
	NT2RM2000566	F-NT2RM2000566	0.39
	NT2RM2000567	F-NT2RM2000567	0.44
25	NT2RM2000569	F-NT2RM2000569	0.49
	NT2RM2000577	F-NT2RM2000577	0.45
	NT2RM2000581	F-NT2RM2000581	0. 72
	NT2RM2000588	F-NT2RM2000588	0. 31
30	NT2RM2000594	F-NT2RM2000594	0.79
	NT2RM2000599	F-NT2RM2000599	0.80
	NT2RM2000612	F-NT2RM2000612	0. 75
	NT2RM2000623	F-NT2RM2000623	0.72
35	NT2RM2000624	F-NT2RM2000624	0. 94
	NT2RM2000635	F-NT2RM2000635	0.73
	NT2RM2000636	F-NT2RM2000636	0. 45
	NT2RM2000639	F-NT2RM2000639	0. 43
40	NT2RM2000649	F-NT2RM2000649	0. 74
10	NT2RM2000669	F-NT2RM2000669	0. 94
	NT2RM2000691	F-NT2RM2000691	0. 84
	NT2RM2000714	F-NT2RM2000714	0. 92
45	NT2RM2000718	F-NT2RM2000718	0. 52
45	NT2RM2000735	F-NT2RM2000735	0. 33
	NT2RM2000740	F-NT2RM2000740	0. 94
	NT2RM2000795	F-NT2RM2000795	0.44
	NT2RM2000821	F-NT2RM2000821	0. 33
50	NT2RM2000837	F-NT2RM2000837	0.57
	NT2RM2000951	F-NT2RM2000951	0. 32
	NT2RM2000952	F-NT2RM2000952	0.50
	NT2RM2000984	F-NT2RM2000984	0.76
55	NT2RM2001004	F-NT2RM2001004	0.55
	NT2RM2001035	F-NT2RM2001035	0. 82

	NT2RM2001065	F-NT2RM2001065	0.65
	NT2RM2001100	F-NT2RM2001100	0.89
5	NT2RM2001105	F-NT2RM2001105	0.94
•	NT2RM2001131	F-NT2RM2001131_	0.94
	NT2RM2001141	F-NT2RM2001141	0.74
	NT2RM2001152	F-NT2RM2001152	0.33
	NT2RM2001177	F-NT2RM2001177	0.32
10	NT2RM2001194	F-NT2RM2001194	0. 83
	NT2RM2001196	F-NT2RM2001196	0.42
	NT2RM2001201	F-NT2RM2001201	0.90
	NT2RM2001221	F-NT2RM2001221	0.39
15	NT2RM2001238	F-NT2RM2001238	0.94
	NT2RM2001243	F-NT2RM2001243	0.44
	NT2RM2001247	F-NT2RM2001247,	0.42
	NT2RM2001256	F-NT2RM2001256	0.60
20	NT2RM2001291	F-NT2RM2001291	0.39
	NT2RM2001306	F-NT2RM2001306	0, 81
	NT2RM2001312	F-NT2RM2001312	0.64
	NT2RM2001319	F-NT2RM2001319	0.53
25	NT2RM2001324	F-NT2RM2001324	0.94
	NT2RM2001345	F-NT2RM2001345	0.94
	NT2RM2001360	F-NT2RM2001360	0.69
	NT2RM2001370	F-NT2RM2001370	0.44
30	NT2RM2001393	F-NT2RM2001393	0.94
	NT2RM2001420	F-NT2RM2001420	0.46
	NT2RM2001424	F-NT2RM2001424	0.94
	NT2RM2001499	F-NT2RM2001499	0.63
35	NT2RM2001504	F-NT2RM2001504	0. 94
	NT2RM2001524	F-NT2RM2001524	0.76
	NT2RM2001544	F-NT2RM2001544	0. 59
	NT2RM2001547	F-NT2RM2001547	0.48
40	NT2RM2001575	F-NT2RM2001575	0. 91
<del></del>	NT2RM2001582	F-NT2RM2001582	0.34
	NT2RM2001588	F-NT2RM2001588	0.48
	NT2RM2001592	F-NT2RM2001592	0.64
	NT2RM2001605	F-NT2RM2001605	0.47
45	NT2RM2001613	F-NT2RM2001613	0.61
	NT2RM2001632	F-NT2RM2001632	0.31
	NT2RM2001635	F-NT2RM2001635	0.39
	NT2RM2001637	F-NT2RM2001637	0.71
50	NT2RM2001641	F-NT2RM2001641	0. 73
	NT2RM2001648	F-NT2RM2001648	0.44
	NT2RM2001652	F-NT2RM2001652	0.40
	NT2RM2001659	F-NT2RM2001659	0.35
55	NT2RM2001664	F-NT2RM2001664	0.53
	NT2RM2001668	F-NT2RM2001668	0.64

	NT2RM2001670	F-NT2RM2001670	0, 59
	NT2RM2001671	F-NT2RM2001671	0. 41
	NT2RM2001675	F-NT2RM2001675	0. 51
5	NT2RM2001681	F-NT2RM2001681	0. 33
	NT2RM2001688	F-NT2RM2001688	0. 74
	NT2RM2001695	F-NT2RM2001695	0, 31
	NT2RM2001696	F-NT2RM2001696	0.94
10	NT2RM2001698	F-NT2RM2001698	0. 69
	NT2RM2001699	F-NT2RM2001699	0. 85
	NT2RM2001700	F-NT2RM2001700	0.71
	NT2RM2001706	F-NT2RM2001706	0.38
15	NT2RM2001716	F-NT2RM2001716	0. 78
	NT2RM2001718	F-NT2RM2001718	0.81
	NT2RM2001723	F-NT2RM2001723	
	NT2RM2001727	F-NT2RM2001727	0. 87
20	NT2RM2001730	F-NT2RM2001730	0.94
	NT2RM2001743	F-NT2RM2001743	0.94
	NT2RM2001753	F-NT2RM2001753	0. 75
	NT2RM2001760	F-NT2RM2001760	0. 75
25	NT2RM2001768	F-NT2RM2001768	0.71
	NT2RM2001771	F-NT2RM2001771	0.33
	NT2RM2001782	F-NT2RM2001782	0.34
	NT2RM2001784	F-NT2RM2001784	0.45
	NT2RM2001785	F-NT2RM2001785	0.94
30	NT2RM2001797	F-NT2RM2001797	0.74
	NT2RM2001800	F-NT2RM2001800	0.90
	NT2RM2001803	F-NT2RM2001803	0.36
	NT2RM2001805	F-NT2RM2001805	0.81
35	NT2RM2001813	F-NT2RM2001813	0.94
	NT2RM2001823	F-NT2RM2001823	0.94
	NT2RM2001839	F-NT2RM2001839	0.39
	NT2RM2001840	F-NT2RM2001840	0.94
40	NT2RM2001855	F-NT2RM2001855	0.92
	NT2RM2001867	F-NT2RM2001867	0. 82
	NT2RM2001879	F-NT2RM2001879	0.86
	NT2RM2001886	F-NT2RM2001886	0.91
45	NT2RM2001896	F-NT2RM2001896	0. 94
	NT2RM2001903	F-NT2RM2001903	0. 61
	NT2RM2001930	F-NT2RM2001930	0. 94
	NT2RM2001935	F-NT2RM2001935	0. 77
50	NT2RM2001936	F-NT2RM2001936	0. 94
	NT2RM2001950	F-NT2RM2001950	0.34
	NT2RM2001982	F-NT2RM2001982	0. 48
	NT2RM2001983	F-NT2RM2001983	0. 59
55	NT2RM2001989	F-NT2RM2001989	0. 94
	NT2RM2001997	F-NT2RM2001997	0. 90

		NT2RM2001998	F-NT2RM2001998	0.72
		NT2RM2002004	F-NT2RM2002004	0. 46
	5	NT2RM2002014	F-NT2RM2002014	0.32
	5	NT2RM2002030	F-NT2RM2002030	0.54
		NT2RM2002049	F-NT2RM2002049	0.59
		NT2RM2002055	F-NT2RM2002055	0.94
	-	NT2RM2002088	F-NT2RM2002088	0. 47
	10	NT2RM2002091	F-NT2RM2002091	0.78
		NT2RM2002100	F-NT2RM2002100	0. 87
		NT2RM2002109	F-NT2RM2002109	0.86
		NT2RM2002128	F-NT2RM2002128	0.42
	15	NT2RM2002142	F-NT2RM2002142	0.94
		NT2RM2002145	F-NT2RM2002145	0.94
		NT2RM2002178	F-NT2RM2002178;	0.44
		NT2RM2002580	F-NT2RM2002580	0.94
	20	NT2RM4000024	F-NT2RM4000024	0.76
		NT2RM4000027	F-NT2RM4000027	0.40
		NT2RM4000030	F-NT2RM4000030	0.66
		NT2RM4000061	F-NT2RM4000061	0. 33
	25	NT2RM4000104	F-NT2RM4000104	0.93
		NT2RM4000139	F-NT2RM4000139	0.87
		NT2RM4000155	F-NT2RM4000155	0.51
		NT2RM4000156	F-NT2RM4000156	0.39
;	o	NT2RM4000167	F-NT2RM4000167	0. 50
		NT2RM4000169	F-NT2RM4000169	0. 44
		NT2RM4000191	F-NT2RM4000191	0.94
		NT2RM4000197	F-NT2RM4000197	0. 41
;	5	NT2RM4000199	F-NT2RM4000199	0. 76
		NT2RM4000202	F-NT2RM4000202	0.47
		NT2RM4000210	F-NT2RM4000210	0. 44
		NT2RM4000215	F-NT2RM4000215	0. 94
4	o	NT2RM4000229	F-NT2RM4000229	0. 93
•		NT2RM4000233	F-NT2RM4000233	0.74
		NT2RM4000251	F-NT2RM4000251	0. 48
		NT2RM4000290	F-NT2RM4000290	0.94
	5	NT2RM4000324		0. 32
•	3	NT2RM4000344	F-NT2RM4000344	0.94
		NT2RM4000349	F-NT2RM4000349	0.78
		NT2RM4000354	F-NT2RM4000354	0. 36
50	_	NT2RM4000356	F-NT2RM4000356	0.58
	U	NT2RM4000386	F-NT2RM4000386	0. 90
		NT2RM4000395	F-NT2RM4000395	0.47
		NT2RM4000421	F-NT2RM4000421	0.46
		NT2RM4000433	F-NT2RM4000433	0.48
5	5	NT2RM4000457	F-NT2RM4000457	0. 75
		NT2RM4000471	F-NT2RM4000471	0. 92

	NT2RM4000486	F-NT2RM4000486	0. 55
	NT2RM4000496	F-NT2RM4000496	0. 62
5	NT2RM4000511	F-NT2RM4000511	0.74
·	NT2RM4000514	F-NT2RM4000514_	0.72
	NT2RM4000515	F-NT2RM4000515	0.66
	NT2RM4000520	F-NT2RM4000520	0.69
	NT2RM4000531	F-NT2RM4000531	0, 62
10	NT2RM4000534	F-NT2RM4000534	0.94
	NT2RM4000585	F-NT2RM4000585	0.52
	NT2RM4000590	F-NT2RM4000590	0.94
	NT2RM4000595	F-NT2RM4000595	0. 44
15	NT2RM4000603	F-NT2RM4000603	0.94
	NT2RM4000611	F-NT2RM4000611	0.46
	NT2RM4000616	F-NT2RM4000616,	0.94
	NT2RM4000674	F-NT2RM4000674	0.80
20	NT2RM4000689	F-NT2RM4000689	0. 58
	NT2RM4000698	F-NT2RM4000698	0.71
	NT2RM4000700	F-NT2RM4000700	0.59
	NT2RM4000712	F-NT2RM4000712	0.94
25	NT2RM4000717	F-NT2RM4000717	0.69
	NT2RM4000733	F-NT2RM4000733	0.94
	NT2RM4000734	F-NT2RM4000734	0.42
•	NT2RM4000741	F-NT2RM4000741	0.83
30	NT2RM4000751	F-NT2RM4000751	0.91
	NT2RM4000764	F-NT2RM4000764	0.35
	NT2RM4000778	F-NT2RM4000778	0.90
	NT2RM4000787	F-NT2RM4000787	0.40
35	NT2RM4000790	F-NT2RM4000790	0.62
	NT2RM4000795	F-NT2RM4000795	0.94
	NT2RM4000796	F-NT2RM4000796	0.93
	NT2RM4000798	F-NT2RM4000798	0.61
40	NT2RM4000813	F-NT2RM4000813	0.58
40	NT2RM4000820	F-NT2RM4000820	0.39
	NT2RM4000833	F-NT2RM4000833	0.80
	NT2RM4000848	F-NT2RM4000848	0.73
	NT2RM4000852	F-NT2RM4000852	0.74
45	NT2RM4000887	F-NT2RM4000887	0. 42
	NT2RM4000895	F-NT2RM4000895	0.47
	NT2RM4000950	F-NT2RM4000950	0. 53
	NT2RM4000971	F-NT2RM4000971	0.73
50	NT2RM4000996	F-NT2RM4000996	0.42
	NT2RM4001002	F-NT2RM4001002	0.57
	NT2RM4001032	F-NT2RM4001032	0.94
	NT2RM4001047	F-NT2RM4001047	0.76
55	NT2RM4001054	F-NT2RM4001054	0. 91
	NT2RM4001084	F-NT2RM4001084	0. 94

	NT2RM4001092	F-NT2RM4001092	0.94
	NT2RM4001116	F-NT2RM4001116	0.36
	NT2RM4001140	F-NT2RM4001140	0.82
5	NT2RM4001151	F-NT2RM4001151	0.94
	NT2RM4001155	F-NT2RM4001155	0. 94
	NT2RM4001187	F-NT2RM4001187	0. 61
	NT2RM4001191	F-NT2RM4001191	0.77
10	NT2RM4001200	F-NT2RM4001200	0.72
	NT2RM4001203	F-NT2RM4001203	0. 60
	NT2RM4001204	F-NT2RM4001204	0.89
		F-NT2RM4001217	0. 93
15	NT2RM4001217		
15	NT2RM4001256	F-NT2RM4001256	0.89
	NT2RM4001258	F-NT2RM4001258	0.94
	NT2RM4001309	F-NT2RM4001309;	
	NT2RM4001316	F-NT2RM4001316	0.38
20	NT2RM4001320	F-NT2RM4001320	0. 33
	NT2RM4001340	F-NT2RM4001340	0, 83
	NT2RM4001344	F-NT2RM4001344	0.94
	NT2RM4001347	F-NT2RM4001347	0.88
25	NT2RM4001371	F-NT2RM4001371	0.61
	NT2RM4001382	F-NT2RM4001382	0. 93
	NT2RM4001384	F-NT2RM4001384	0. 32
	NT2RM4001410	F-NT2RM4001410	0.94
30	NT2RM4001411	F-NT2RM4001411	0. 47
30	NT2RM4001412	F-NT2RM4001412	0.85
	NT2RM4001444	F-NT2RM4001444	0. 94
	NT2RM4001454	F-NT2RM4001454	0. 72
	NT2RM4001455	F-NT2RM4001455	0.39
35	NT2RM4001483	F-NT2RM4001483	0.76
	NT2RM4001489	F-NT2RM4001489	0. 94
	NT2RM4001522	F-NT2RM4001522	0.88
	NT2RM4001565	F-NT2RM4001565	0.78
40	NT2RM4001566	F-NT2RM4001566	0.87
	NT2RM4001569	F-NT2RM4001569	0.70
	NT2RM4001582	F-NT2RM4001582	0.76
	NT2RM4001592	F-NT2RM4001592	0.31
45	NT2RM4001594	F-NT2RM4001594	0.86
-	NT2RM4001597	F-NT2RM4001597	0. 91
	NT2RM4001611	F-NT2RM4001611	0.91
	NT2RM4001629	F-NT2RM4001629	0.72
	NT2RM4001650	F-NT2RM4001650	0. 37
50	NT2RM4001662	F-NT2RM4001662	0.70
	NT2RM4001666	F-NT2RM4001666	0.37
	NT2RM4001682	F-NT2RM4001682	0. 91
	NT2RM4001710	F-NT2RM4001710	0. 45
55	NT2RM4001714	F-NT2RM4001714	0. 94
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	NT2RM4001715	F-NT2RM4001715	0. 33
	NT2RM4001731	F-NT2RM4001731	0. 94
	NT2RM4001741	F-NT2RM4001741	0. 87
5	NT2RM4001746	F-NT2RM4001746_	
	NT2RM4001758	F-NT2RM4001758	0. 59
	NT2RM4001783	F-NT2RM4001783	0.68
	NT2RM4001813	F-NT2RM4001813	0.39
10	NT2RM4001819	F-NT2RM4001819	0. 43
	NT2RM4001823	F-NT2RM4001823	0. 94
	NT2RM4001828	F-NT2RM4001828	0. 83
	NT2RM4001836	F-NT2RM4001836	0. 61
15	NT2RM4001841	F-NT2RM4001841	0. 37
	NT2RM4001842	F-NT2RM4001842	0. 68
	NT2RM4001865	F-NT2RM4001865	
	NT2RM4001876	F-NT2RM4001876	0.88
20	NT2RM4001880	F-NT2RM4001880	0. 43
	NT2RM4001905	F-NT2RM4001905	0.94
	NT2RM4001922	F-NT2RM4001922	0.48
	NT2RM4001938	F-NT2RM4001938	0. 67
25	NT2RM4001940	F-NT2RM4001940	0. 92
25	NT2RM4001965	F-NT2RM4001965	0.94
	NT2RM4001969	F-NT2RM4001969	0.88
	NT2RM4001979	F-NT2RM4001979	0.42
20	NT2RM4001987	F-NT2RM4001987	0.86
30	NT2RM4002013	F-NT2RM4002013	0.94
	NT2RM4002018	F-NT2RM4002018	0. 92
	NT2RM4002034	F-NT2RM4002034	0. 43
	NT2RM4002044	F-NT2RM4002044	0. 42
35	NT2RM4002054	F-NT2RM4002054	0.88
	NT2RM4002055	F-NT2RM4002055	0. 32
	NT2RM4002062	F-NT2RM4002062	0. 87
	NT2RM4002063	F-NT2RM4002063	0.49
40	NT2RM4002066	F-NT2RM4002066	0. 39
	NT2RM4002073	F-NT2RM4002073	0. 78
	NT2RM4002075		
	NT2RM4002093		0. 82
45	NT2RM4002109	F-NT2RM4002109	0. 52
	NT2RM4002128	F-NT2RM4002128	0. 68
	NT2RM4002140	F-NT2RM4002140	0.94
	NT2RM4002145	F-NT2RM4002145	0. 94
50	NT2RM4002146	F-NT2RM4002146	0. 89
	NT2RM4002161	F-NT2RM4002161	0. 49
	NT2RM4002174	F-NT2RM4002174	0. 36
	NT2RM4002189	F-NT2RM4002189	0. 34
55	NT2RM4002194	F-NT2RM4002194	0. 73
	NT2RM4002205	F-NT2RM4002205	0. 94

	NT2RM4002213	F-NT2RM4002213	0. 52
	NT2RM4002226	F-NT2RM4002226	0. 48
	NT2RM4002251	F-NT2RM4002251	0. 94
5	NT2RM4002256	F-NT2RM4002256_	
			-
	NT2RM4002266	F-NT2RM4002266	0.73
	NT2RM4002294	F-NT2RM4002294	0.86
10	NT2RM4002301	F-NT2RM4002301	0.40
	NT2RM4002323	F-NT2RM4002323	0.81
	NT2RM4002339	F-NT2RM4002339	0.32
	NT2RM4002344	F-NT2RM4002344	0. 87
15	NT2RM4002373	F-NT2RM4002373	0.41
73	NT2RM4002374	F-NT2RM4002374	0. 33
	NT2RM4002398	F-NT2RM4002398	0.94
	NT2RM4002409	F-NT2RM4002409,	
·	NT2RM4002446	F-NT2RM4002446	0.61
20	NT2RM4002452	F-NT2RM4002452	0. 47
	NT2RM4002457	F-NT2RM4002457	0. 92
	NT2RM4002460	F-NT2RM4002460	0.61
	NT2RM4002482	F-NT2RM4002482	0.94
25	NT2RM4002493	F-NT2RM4002493	0. 39
	NT2RM4002527	F-NT2RM4002527	0.64
	NT2RM4002534	F-NT2RM4002534	0. 76
	NT2RM4002558	F-NT2RM4002558	0.62
30	NT2RM4002565	F-NT2RM4002565	0. 54
	NT2RM4002571	F-NT2RM4002571	0. 90
	NT2RM4002593	F-NT2RM4002593	0. 91
	NT2RM4002594	F-NT2RM4002594	0. 70
35	NT2RM4002623	F-NT2RM4002623	0. 81
-	NT2RP1000018	F-NT2RP1000018	0.31
	NT2RP1000035	F-NT2RP1000035	0.49
	NT2RP1000040	F-NT2RP1000040	0. 45
40	NT2RP1000063	F-NT2RP1000063	0.90
40	NT2RP1000086	F-NT2RP1000086	0.89
	NT2RP1000101	F-NT2RP1000101	0. 76
	NT2RP1000111	F-NT2RP1000111	0. 94
	NT2RP1000112	F-NT2RP1000112	0. 45
45	NT2RP1000124	F-NT2RP1000124	0. 94
	NT2RP1000130	F-NT2RP1000130	0. 69
	NT2RP1000163	F-NT2RP1000163	0. 90
	NT2RP1000170	F-NT2RP1000170	0. 94
50	NT2RP1000174	F-NT2RP1000174	0. 74
	NT2RP1000191	F-NT2RP1000191	0. 76
	NT2RP1000202	F-NT2RP1000202	0.60
	NT2RP1000243	F-NT2RP1000243	0. 81
55	NT2RP1000259	F-NT2RP1000259	0. 79
	NT2RP1000272	F-NT2RP1000272	0. 94

	NT2RP1000324	F-NT2RP1000324	0. 32
	NT2RP1000326	F-NT2RP1000326	0. 94
	NT2RP1000333	F-NT2RP1000333	0.94
5	NT2RP1000348	F-NT2RP1000348	0.94
	NT2RP1000357	F-NT2RP1000357	0.87
	NT2RP1000358	F-NT2RP1000358	0.79
	NT2RP1000363	F-NT2RP1000363	0.40
10	NT2RP1000376	F-NT2RP1000376	0.61
	NT2RP1000409	F-NT2RP1000409	0.85
	NT2RP1000413	F-NT2RP1000413	0. 42
	NT2RP1000416	F-NT2RP1000416	0. 68
15	NT2RP1000418	F-NT2RP1000418	0.88
,,,	NT2RP1000418		
		F-NT2RP1000439	0.69
	NT2RP1000443	F-NT2RP1000443	
	NT2RP1000460	F-NT2RP1000460	0.86
20	NT2RP1000470	F-NT2RP1000470	0.56
	NT2RP1000478	F-NT2RP1000478	0.87
	NT2RP1000481	F-NT2RP1000481	0. 58
	NT2RP1000493	F-NT2RP1000493	0. 34
25	NT2RP1000513	F-NT2RP1000513	0.89
	NT2RP1000522	F-NT2RP1000522	0.34
	NT2RP1000547	F-NT2RP1000547	0. 89
	NT2RP1000574	F-NT2RP1000574	0.74
30	NT2RP1000577	F-NT2RP1000577	0. 53
50	NT2RP1000581	F-NT2RP1000581	0.66
	NT2RP1000609	F-NT2RP1000609	0. 90
	NT2RP1000629	F-NT2RP1000629	0.94
	NT2RP1000630	F-NT2RP1000630	0.93
35	NT2RP1000677	F-NT2RP1000677	0. 94
	NT2RP1000688	F-NT2RP1000688	0. 94
	NT2RP1000695	F-NT2RP1000695	0. 89
	NT2RP1000701	F-NT2RP1000701	0.89
40	NT2RP1000721	F-NT2RP1000721	0. 91
	NT2RP1000730	F-NT2RP1000730	0. 94
	NT2RP1000733	F-NT2RP1000733	
	NT2RP1000738		0.64
45	NT2RP1000746	F-NT2RP1000746	0. 92
,,	NT2RP1000767	F-NT2RP1000767	0. 82
	NT2RP1000782	F-NT2RP1000782	0. 94
	NT2RP1000796	F-NT2RP1000796	0. 33
	NT2R11000825	F-NT2RP1000825	0. 74
50	NT2RT 1000823	F-NT2RP1000833	0. 74
	NT2RP1000834	F-NT2RP1000834	0.88
	NT2RP1000836	F-NT2RP1000836	0.59
55	NT2RP1000846	F~NT2RP1000846	0.46
	NT2RP1000851	F-NT2RP1000851	0. 82

		NT2RP1000856	F-NT2RP1000856	0.94
		NT2RP1000860	F-NT2RP1000860	0.60
		NT2RP1000902	F-NT2RP1000902	0.42
5		NT2RP1000915	F-NT2RP1000915_	0.90
		NT2RP1000916	F-NT2RP1000916	0. 67
		NT2RP1000943	F-NT2RP1000943	0.77
		NT2RP1000944	F-NT2RP1000944	0.94
10	•	NT2RP1000947	F-NT2RP1000947	0.38
		NT2RP1000954	F-NT2RP1000954	0.59
		NT2RP1000958	F-NT2RP1000958	0. 59
		NT2RP1000959	F-NT2RP1000959	0.74
15		NT2RP1000966	F-NT2RP1000966	0.93
		NT2RP1000980	F-NT2RP1000980	0.31
		NT2RP1000988	F-NT2RP1000988	0.53
		NT2RP1001011	F-NT2RP1001011	0.65
20		NT2RP1001013	F-NT2RP1001013	0.34
		NT2RP1001014	F-NT2RP1001014	0.94
		NT2RP1001033	F-NT2RP1001033	0.40
		NT2RP1001073	F-NT2RP1001073	0.74
25		NT2RP1001079	F-NT2RP1001079	0.48
		NT2RP1001080	F-NT2RP1001080	0. 91
		NT2RP1001113	F-NT2RP1001113	0.94
		NT2RP1001173	F-NT2RP1001173	0. 57
30		NT2RP1001177	F-NT2RP1001177	0.75
		NT2RP1001185	F-NT2RP1001185	0.94
		NT2RP1001199	F-NT2RP1001199	0. 42
		NT2RP1001247	F-NT2RP1001247	0.67
35		NT2RP1001248	F-NT2RP1001248	0.89
		NT2RP1001253 NT2RP1001286	F-NT2RP1001253 F-NT2RP1001286	0. 94 0. 59
		NT2RP1001294	F-NT2RP1001294	0. 59
		NT2RP1001302	F-NT2RP1001302	0.86
40		NT2RP1001310	F-NT2RP1001310	0. 91
		NT2RP1001311	F-NT2RP1001311	0. 87
		NT2RP1001313	F-NT2RP1001313	0. 91
		NT2RP1001361	F-NT2RP1001361	0.83
45		NT2RP1001385	F-NT2RP1001385	0.36
		NT2RP1001395	F-NT2RP1001395	0.84
		NT2RP1001410	F-NT2RP1001410	0.72
		NT2RP1001424	F-NT2RP1001424	0. 81
50		NT2RP1001432	F-NT2RP1001432	0.81
		NT2RP1001449	F-NT2RP1001449	0.77
		NT2RP1001457	F-NT2RP1001457	0.92
		NT2RP1001466	F-NT2RP1001466	0.44
55		NT2RP1001475	F-NT2RP1001475	0.53
		NT2RP1001482	F-NT2RP1001482	0.34

	NT2RP1001494	F-NF2RP1001494	0. 93
	NT2RP1001543	F-NT2RP1001543	0. 94
	NT2RP1001546	F-NT2RP1001546	0.94
5	NT2RP1001569	F-NT2RP1001569_	
	NT2RP1001616	F-NT2RP1001616	0. 46
	NT2RP1001665	F-NT2RP1001665	0.94
	NT2RP2000001	F-NT2RP2000001	0.94
10	NT2RP2000006	F-NT2RP2000006	0.74
	NT2RP2000007	F-NT2RP2000007	0.94
	NT2RP2000008	F-NT2RP2000008	0.83
	NT2RP2000032	F-NT2RP2000032	0.94
15	NT2RP2000040	F-NT2RP2000040	0. 92
	NT2RP2000045	F-NT2RP2000045	0.94
	NT2RP2000054	F-NT2RP2000054,	0. 53
	NT2RP2000056	F-NT2RP2000056	0. 42
20	NT2RP2000067	F-NT2RP2000067	0.88
	NT2RP2000070	F-NT2RP2000070	0.48
	NT2RP2000079	F-NT2RP2000079	0.85
	NT2RP2000088	F-NT2RP2000088	0.94
25	NT2RP2000091	F-NT2RP2000091	0.79
	NT2RP2000097	F-NT2RP2000097	0.66
	NT2RP2000114	F-NT2RP2000114	0.86
	NT2RP2000120	F-NT2RP2000120	0.48
30	NT2RP2000126	F-NT2RP2000126	0.77
•	NT2RP2000133	F-NT2RP2000133	0.64
	NT2RP2000147	F-NT2RP2000147	0.94
	NT2RP2000153	F-NT2RP2000153	0. 87
35	NT2RP2000157	F-NT2RP2000157	0.48
	NT2RP2000161	F-NT2RP2000161	0.79
	NT2RP2000173	F-NT2RP2000173	0. 75
	NT2RP2000175	F-NT2RP2000175	0.79
40	NT2RP2000195	F-NT2RP2000195	0.91
70	NT2RP2000205	F-NT2RP2000205	0.94
	NT2RP2000208	F-NT2RP2000208	0. 84
	NT2RP2000224	F-NT2RP2000224	0. 52
45	NT2RP2000232		0. 33
45	NT2RP2000233	F-NT2RP2000233	0. 57
	NT2RP2000239	F-NT2RP2000239	0. 52
	NT2RP2000248	F-NT2RP2000248	0. 36
	NT2RP2000274	F-NT2RP2000274	0. 55
50	NT2RP2000283	F-NT2RP2000283	0. 61
	NT2RP2000288	F-NT2RP2000288	0. 31
	NT2RP2000297	F-NT2RP2000297	0. 78
	NT2RP2000298	F-NT2RP2000298	0. 94
55	NT2RP2000310	F-NT2RP2000310	0. 52
	NT2RP2000328	F-NT2RP2000328	0. 75

	NT2RP2000329	F-NT2RP2000329	0. 94
	NT2RP2000346	F-NT2RP2000346	0. 62
	NT2RP2000369	F-NT2RP2000369	0. 87
5	NT2RP2000412	F-NT2RP2000412	
	NT2RP2000414	F-NT2RP2000414	0. 79
	NT2RP2000422	F-NT2RP2000422	0.46
	NT2RP2000438	F-NT2RP2000438	0. 94
10	NT2RP2000448	F-NT2RP2000448	0.94
	NT2RP2000503	F-NT2RP2000503	0.94
	NT2RP2000510	F-NT2RP2000510	0.87
	NT2RP2000516	F-NT2RP2000516	0.42
15	NT2RP2000617	F-NT2RP2000617	0.94
	NT2RP2000634	F-NT2RP2000634	0.74
	NT2RP2000656	F-NT2RP2000656,	0.68
	NT2RP2000658	F-NT2RP2000658	0. 55
20	NT2RP2000668	F-NT2RP2000668	0.81
	NT2RP2000704	F-NT2RP2000704	0.88
	NT2RP2000710	F-NT2RP2000710	0.58
	NT2RP2000764	F-NT2RP2000764	0.59
25	NT2RP2000809	F-NT2RP2000809	0.51
	NT2RP2000812	F-NT2RP2000812	0.37
	NT2RP2000814	F-NT2RP2000814	0.66
	NT2RP2000816	F-NT2RP2000816	0.50
30	NT2RP2000819	F-NT2RP2000819	0. 35
	NT2RP2000841	F-NT2RP2000841	0. 93
	NT2RP2000845	F-NT2RP2000845	0.78
	NT2RP2000863	F-NT2RP2000863	0. 41
35	NT2RP2000880	F-NT2RP2000880	0. 79
	NT2RP2000892	F-NT2RP2000892	0.94
	NT2RP2000931	F-NT2RP2000931	0.89
	NT2RP2000932	F-NT2RP2000932 F-NT2RP2000938	0. 94
40	NT2RP2000938 NT2RP2000943	F-NT2RP2000943	0. 90 0. 61
	NT2RP2000943	F-NT2RP2000945	0.80
	NT2RP2000985	F-NT2RP2000985	0. 94
	NT2RP2001036	F-NT2RP2001036	0.86
45	NT2RP2001044	F-NT2RP2001044	0. 93
	NT2RP2001056	F-NT2RP2001056	0. 77
	NT2RP2001065	F-NT2RP2001065	0. 73
	NT2RP2001070	F-NT2RP2001070	0. 91
50	NT2RP2001081	F-NT2RP2001081	0. 94
	NT2RP2001094	F-NT2RP2001094	0. 93
	NT2RP2001119	F-NT2RP2001119	0. 31
	NT2RP2001127	F-NT2RP2001127	0. 41
55	NT2RP2001137	F-NT2RP2001137	0. 93
55	NT2RP2001168	F-NT2RP2001168	0.71

	NT2RP2001173	F-NT2RP2001173	0.94
	NT2RP2001174	F-NT2RP2001174	0. 71
_	NT2RP2001218	F-NT2RP2001218	0.76
5	NT2RP2001233	F-NT2RP2001233	0.31
	NT2RP2001245	F-NT2RP2001245	0. 33
	NT2RP2001268	F-NT2RP2001268	0. 61
	NT2RP2001327	F-NT2RP2001327	0.86
10	NT2RP2001328	F-NT2RP2001328	0.94
	NT2RP2001366	F-NT2RP2001366	0. 52
	NT2RP2001378	F-NT2RP2001378	0.64
	NT2RP2001381	F-NT2RP2001381	0.80
15	NT2RP2001392	F-NT2RP2001392	0.69
	NT2RP2001394	F-NT2RP2001394	0.38
	NT2RP2001397	F-NT2RP2001397	
	NT2RP2001420	F-NT2RP2001420	0. 88
20	NT2RP2001427	F-NT2RP2001427	0.73
	NT2RP2001436	F-NT2RP2001436	0. 32
	NT2RP2001440	F-NT2RP2001440	0.94
	NT2RP2001450	F-NT2RP2001450	0.38
25	NT2RP2001511	F-NT2RP2001511	0. 53
	NT2RP2001520	F-NT2RP2001520	0.88
	NT2RP2001536	F-NT2RP2001536	0.64
	NT2RP2001560	F-NT2RP2001560	0.79
30	NT2RP2001576	F-NT2RP2001576	0.62
	NT2RP2001581	F-NT2RP2001581	0.48
	NT2RP2001597	F-NT2RP2001597	0.94
	NT2RP2001601	F-NT2RP2001601	0. 33
35	NT2RP2001613	F-NT2RP2001613	0. 41
	NT2RP2001628	F-NT2RP2001628	0. 79
	NT2RP2001634	F-NT2RP2001634	0.87
	NT2RP2001660	F-NT2RP2001660	0.86
40	NT2RP2001675	F-NT2RP2001675	0. 90
40	NT2RP2001721	F-NT2RP2001721	0.32
	NT2RP2001748	F-NT2RP2001748	0.32
		F-NT2RP2001839	0.83
	NT2RP2001861	F-NT2RP2001861	0.33
45	NT2RP2001876	F-NT2RP2001876	0.94
	NT2RP2001883 NT2RP2001898	F-NT2RP2001883	0.90
		F-NT2RP2001898	0.94
	NT2RP2001900	F-NT2RP2001900	0. 93
50	NT2RP2001907 NT2RP2001946	F-NT2RP2001907	0.94
	NT2RP2001946 NT2RP2001947	F-NT2RP2001946	0.63
	NT2RP2001947	F-NT2RP2001947 F-NT2RP2001969	0. 75 0. 94
	NT2RP2001969	F-NT2RP2001969	0. 48
55	NT2RP2001976	F-NT2RP2001976	0. 44
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	NT2RP2001991	F-NT2RP2001991	0. 39
	NT2RP2002025	F-NT2RP2002025	0.94
5	NT2RP2002033	F-NT2RP2002033	0.40
5	NT2RP2002046	F-NT2RP2002046	0, 56
	NT2RP2002058	F-NT2RP2002058	0.64
	NT2RP2002076	F-NT2RP2002076	0.40
	NT2RP2002078	F-NT2RP2002078	0. 81
10	NT2RP2002079	F-NT2RP2002079	0. 82
	NT2RP2002099	F-NT2RP2002099	0. 92
	NT2RP2002154	F-NT2RP2002154	0. 44
	NT2RP2002185	F-NT2RP2002185	0.94
15	NT2RP2002193	F-NT2RP2002193	0. 48
	NT2RP2002208	F-NT2RP2002208	0.49
	NT2RP2002231	F-NT2RP2002231	0. 61
	NT2RP2002235	F-NT2RP2002235	0.87
20	NT2RP2002252	F-NT2RP2002252	0.51
	NT2RP2002270	F-NT2RP2002270	0.78
	NT2RP2002292	F-NT2RP2002292	0.94
	NT2RP2002312	F-NT2RP2002312	0.33
25	NT2RP2002325	F-NT2RP2002325	0.53
	NT2RP2002333	F-NT2RP2002333	0.60
	NT2RP2002373	F-NT2RP2002373	0.91
	NT2RP2002385	F-NT2RP2002385	0.94
30	NT2RP2002408	F-NT2RP2002408	0.44
	NT2RP2002426	F-NT2RP2002426	0.54
	NT2RP2002442	F-NT2RP2002442	0.37
	NT2RP2002464	F-NT2RP2002464	0. 79
35	NT2RP2002479	F-NT2RP2002479	0. 92
	NT2RP2002498	F-NT2RP2002498	0.94
	NT2RP2002503	F-NT2RP2002503	0. 90
	NT2RP2002520	F-NT2RP2002520	0.37
40	NT2RP2002537	F-NT2RP2002537	0. 42
40	NT2RP2002549	F-NT2RP2002549	0. 75
	NT2RP2002595	F-NT2RP2002595	0. 90
	NT2RP2002609	F-NT2RP2002609	0.33
	NT2RP2002618	F-NT2RP2002618	0.52
45	NT2RP2002621	F-NT2RP2002621	0.67
	NT2RP2002672	F-NT2RP2002672	0.43
	NT2RP2002701	F-NT2RP2002701	0.69
	NT2RP2002706	F-NT2RP2002706	0.51
50	NT2RP2002710	F-NT2RP2002710	0.94
	NT2RP2002769	F-NT2RP2002769	0.42
	NT2RP2002800	F-NT2RP2002800	0.47
	NT2RP2002862	F-NT2RP2002862	0.88
55	NT2RP2002880	F-NT2RP2002880	0.78
	NT2RP2002891	F-NT2RP2002891	0.69

	NT2RP2002925	F-NT2RP2002925	0. 65
	NT2RP2002928	F-NT2RP2002928	0. 47
	NT2RP2002929	F-NT2RP2002929	0. 94
5	NT2RP2002939	F-NT2RP2002939	0.94
	NT2RP2002954	F-NT2RP2002954	0. 94
	NT2RP2002959	F-NT2RP2002959	0. 45
	NT2RP2002979	F-NT2RP2002979	0.51
10	NT2RP2002980	F-NT2RP2002980	0. 38
	NT2RP2002986	F-NT2RP2002986	0. 34
	NT2RP2002993	F-NT2RP2002993	0. 94
	NT2RP2003034	F-NT2RP2003034	0. 34
15	NT2RP2003099	F-NT2RP2003099	0. 46
	NT2RP2003108	F-NT2RP2003108	0. 67
	NT2RP2003117	F-NT2RP2003117,	0. 57
	NT2RP2003121	F-NT2RP2003121	0.94
20	NT2RP2003125	F-NT2RP2003125	0.94
r	NT2RP2003137	F-NT2RP2003137	0. 60
	NT2RP2003157	F-NT2RP2003157	0.94
	NT2RP2003158	F-NT2RP2003158	0. 75
25	NT2RP2003165	F-NT2RP2003165	0.90
	NT2RP2003177	F-NT2RP2003177	0.85
	NT2RP2003194	F-NT2RP2003194	0.94
	NT2RP2003228	F-NT2RP2003228	0.57
30	NT2RP2003243	F-NT2RP2003243	0.94
30	NT2RP2003265	F-NT2RP2003265	0.68
	NT2RP2003272	F-NT2RP2003272	0.94
	NT2RP2003277	F-NT2RP2003277	0. 73
	NT2RP2003280	F-NT2RP2003280	0.94
35	NT2RP2003286	F-NT2RP2003286	0.44
	NT2RP2003293	F-NT2RP2003293	0. 42
	NT2RP2003295	F-NT2RP2003295	0.89
	NT2RP2003297	F-NT2RP2003297	0.94
40	NT2RP2003307	F-NT2RP2003307	0. 79
	NT2RP2003308	F-NT2RP2003308	0.49
		F-NT2RP2003329	0. 33
	NT2RP2003347	F-NT2RP2003347	0. 79
45	NT2RP2003367	F-NT2RP2003367	0.94
	NT2RP2003391	F-NT2RP2003391	0.90
	NT2RP2003393	F-NT2RP2003393	0.94
	NT2RP2003401	F-NT2RP2003401	0.80
50	NT2RP2003433	F-NT2RP2003433	0.40
	NT2RP2003445	F-NT2RP2003445	0.94
	NT2RP2003446	F-NT2RP2003446	0.40
	NT2RP2003466	F-NT2RP2003466	0. 58
55	NT2RP2003480	F-NT2RP2003480	0.59
	NT2RP2003506	F-NT2RP2003506	0. 94

	NT2RP2003511	F-NT2RP2003511	0: 31
	NT2RP2003513	F-NT2RP2003513	0. 67
_	NT2RP2003533	F-NT2RP2003533	0. 57
5	NT2RP2003543	F-NT2RP2003543	0. 94
	NT2RP2003564	F-NT2RP2003564	0.88
	NT2RP2003567	F-NT2RP2003567	0.69
	NT2RP2003581	F-NT2RP2003581	0.40
10	NT2RP2003596	F-NT2RP2003596	0.83
	NT2RP2003604	F-NT2RP2003604	0. 87
	NT2RP2003629	F-NT2RP2003629	0. 87
	NT2RP2003643	F-NT2RP2003643	0. 94
15	NT2RP2003687	F-NT2RP2003687	0. 31
	NT2RP2003691	F-NT2RP2003691	0.34
	NT2RP2003702	F-NT2RP2003702,	
	NT2RP2003713	F~NT2RP2003713	0. 53
20	NT2RP2003714	F-NT2RP2003714	0. 61
	NT2RP2003737	F-NT2RP2003737	0.88
	NT2RP2003751	F-NT2RP2003751	0. 36
	NT2RP2003760	F-NT2RP2003760	0. 62
25	NT2RP2003764	F-NT2RP2003764	0. 94
	NT2RP2003769	F-NT2RP2003769	0. 69
	NT2RP2003777	F-NT2RP2003777	0. 93
	NT2RP2003781	F-NT2RP2003781	0.37
30	NT2RP2003793	F-NT2RP2003793	0.92
	NT2RP2003825	F-NT2RP2003825	0.49
	NT2RP2003840	F-NT2RP2003840	0.88
	NT2RP2003857	F-NT2RP2003857	0.34
35	NT2RP2003952	F-NT2RP2003952	0.89
00	NT2RP2003976	F-NT2RP2003976	0.33
	NT2RP2003981	F-NT2RP2003981	0.64
	NT2RP2003984	F-NT2RP2003984	0. 41
40	NT2RP2003986	F-NT2RP2003986	0. 32
40	NT2RP2004013	F-NT2RP2004013	0.48
	NT2RP2004041	F-NT2RP2004041	0. 78
	NT2RP2004042	F-NT2RP2004042	0. 66
	NT2RP2004066	F-NT2RP2004066	0. 56
45	NT2RP2004081	F-NT2RP2004081	0.94
	NT2RP2004098	F-NT2RP2004098	0. 39
	NT2RP2004124	F-NT2RP2004124	0. 75
	NT2RP2004152	F-NT2RP2004152	0. 54
50	NT2RP2004165	F-NT2RP2004165	0. 68
	NT2RP2004187	F-NT2RP2004187	0. 92
	NT2RP2004194	F-NT2RP2004194	0. 90
	NT2RP2004196		0. 94
55	NT2RP2004226		0. 80
	NT2RP2004239	F-NT2RP2004239	0. 63

	NT2RP2004240	F-NT2RP2004240	0. 86
	NT2RP2004242	F-NT2RP2004242	0. 94
	NT2RP2004245	F-NT2RP2004245	0. 92
5	NT2RP2004316	F-NT2RP2004316	0. 33
	NT2RP2004364	F-NT2RP2004364	
	NT2RP2004365	F-NT2RP2004365	0. 43
	NT2RP2004366	F-NT2RP2004366	0.32
10	NT2RP2004373	F-NT2RP2004373	0. 60
	NT2RP2004389	F-NT2RP2004389	0. 94
	NT2RP2004392	F-NT2RP2004392	0. 93
	NT2RP2004399	F-NT2RP2004399	0.36
15	NT2RP2004463	F-NT2RP2004463	0.43
	NT2RP2004476	F-NT2RP2004476	0.80
	NT2RP2004538	F-NT2RP2004538,	0. 94
	NT2RP2004551	F-NT2RP2004551	0.47
20	NT2RP2004568	F-NT2RP2004568	0.73
	NT2RP2004594	F-NT2RP2004594	0.64
	NT2RP2004600	F-NT2RP2004600	0. 53
	NT2RP2004602	F-NT2RP2004602	0.33
25	NT2RP2004614	F-NT2RP2004614	0. 34
	NT2RP2004655	F-NT2RP2004655	0.77
	NT2RP2004664	F-NT2RP2004664	0. 94
	NT2RP2004689	F-NT2RP2004689	0.83
30	NT2RP2004710	F-NT2RP2004710	0. 94
	NT2RP2004743		0. 32
	NT2RP2004768		0. 53
	NT2RP2004791	F-NT2RP2004791	0. 64
35	NT2RP2004799	F-NT2RP2004799	
	NT2RP2004802	F-NT2RP2004802	
	NT2RP2004816		0. 78
	NT2RP2004841		0. 61
40	NT2RP2004861		0. 32
	NT2RP2004897		0. 67
	NT2RP2004933	F-NT2RP2004933	0. 94
	NT2RP2004936	1 1112111 200-1300	
45	NT2RP2004959 NT2RP2004978		0. 78
40	NT2RP2004978		0.86
	NT2RP2004985	F-NT2RP2004985	0. 42
	NT2RP2005000	F-NT2RP2004999	0. 55
	NT2RP2005001	F-NT2RP2005000	0.40
50	NT2RP2005003	F-NT2RP2005001	0. 94
	NT2RP2005003	F-NT2RP2005003 F-NT2RP2005012	0. 44 0. 73
•	NT2RP2005012	F-NT2RP2005012	
	NT2RP2005018	F-NT2RP2005018	0. 43 0. 34
55	NT2RP2005037	F-NT2RP2005037	0. 94
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	NT2RP2005038	F-NT2RP2005038	0.56
	NT2RP2005116	F-NT2RP2005116	0.94
	NT2RP2005126	F-NT2RP2005126	0.69
5	NT2RP2005139	F-NT2RP2005139	0.94
	NT2RP2005140	F-NT2RP2005140	0.32
	NT2RP2005144	F-NT2RP2005144	0.39
	NT2RP2005147	F-NT2RP2005147	0. 32
10	NT2RP2005159	F-NT2RP2005159	0. 52
	NT2RP2005162	F-NT2RP2005162	0.86
	NT2RP2005168	F-NT2RP2005168	0. 53
	NT2RP2005204	F-NT2RP2005204	0. 94
15	NT2RP2005227	F-NT2RP2005227	0.53
	NT2RP2005239	F-NT2RP2005239	0. 91
	NT2RP2005270	F-NT2RP2005270,	0.72
	NT2RP2005276	F-NT2RP2005276	0.41
20	NT2RP2005287	F-NT2RP2005287	0.44
	NT2RP2005288	F-NT2RP2005288	0.82
	NT2RP2005315	F-NT2RP2005315	0. 79
	NT2RP2005358	F-NT2RP2005358	0. 94
25	NT2RP2005393	F-NT2RP2005393	0. 41
	NT2RP2005436	F-NT2RP2005436	0. 45
	NT2RP2005441	F-NT2RP2005441	0. 88
	NT2RP2005453	F-NT2RP2005453	0. 59
30	NT2RP2005457	F-NT2RP2005457	0. 56
	NT2RP2005464	F-NT2RP2005464	0. 72
	NT2RP2005465	F-NT2RP2005465	0. 40
	NT2RP2005472	F-NT2RP2005472	0. 42
25	NT2RP2005490	F-NT2RP2005490	0. 52
35	NT2RP2005495	F-NT2RP2005495	0.94
	NT2RP2005496	F-NT2RP2005496	0.52
	NT2RP2005498	F-NT2RP2005498	0.74
	NT2RP2005509	F-NT2RP2005509	0.55
40	NT2RP2005520	F-NT2RP2005520	0. 54
	NT2RP2005525	F-NT2RP2005525	0.41
	NT2RP2005539 NT2RP2005540	F-NT2RP2005539 F-NT2RP2005540	0. 94 0. 32
	NT2RP2005549		0. 32
45	NT2RP2005555	F-NT2RP2005549 F-NT2RP2005555	0. 79
	NT2RP2005557	F-NT2RP2005557	0. 03
	NT2RP2005600	F-NT2RP2005600	0. 76
	NT2RP2005605	F-NT2RP2005605	0. 76
50	NT2RP2005620	F-NT2RP2005620	0. 66
	NT2RP2005620	F-NT2RP2005620	0. 37
	NT2RP2005625	F-NT2RP2005635	0. 37
	NT2RP2005637	F-NT2RP2005637	0. 94
55	NT2RP2005640	F-NT2RP2005640	0. 87
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	NT2RP2005654	F-NT2RP2005654	0.94
	NT2RP2005669	F-NT2RP2005669	0.94
	NT2RP2005675	F-NT2RP2005675	0.72
5	NT2RP2005683	F-NT2RP2005683	0.34
	NT2RP2005690	F-NT2RP2005690	0. 54
	NT2RP2005712	F-NT2RP2005712	0. 93
	NT2RP2005722	F-NT2RP2005722	0.50
10	NT2RP2005723	F-NT2RP2005723	0. 45
	NT2RP2005732	F-NT2RP2005732	0. 54
	NT2RP2005748		0.76
	NT2RP2005752	F-NT2RP2005752	0. 47
15	NT2RP2005763	F-NT2RP2005763	0. 87
	NT2RP2005767		0. 58
	NT2RP2005773	F-NT2RP2005773,	
	NT2RP2005775	•	0.34
20	NT2RP2005781	F-NT2RP2005781	0.81
	NT2RP2005784	F-NT2RP2005784	0.94
	NT2RP2005804	F-NT2RP2005804	0.94
	NT2RP2005812	F-NT2RP2005812	0.90
25	NT2RP2005835	F-NT2RP2005835	0.94
	NT2RP2005853	F-NT2RP2005853	0.77
	NT2RP2005859	F-NT2RP2005859	0.68
	NT2RP2005868	F-NT2RP2005868	0.76
30	NT2RP2005886	F-NT2RP2005886	0.62
30	NT2RP2005890	F-NT2RP2005890	0. 79
	NT2RP2005901	F-NT2RP2005901	0.94
	NT2RP2005933	F-NT2RP2005933	0. 55
	NT2RP2006023	F-NT2RP2006023	0. 47
35	NT2RP2006038	F-NT2RP2006038	0.73
	NT2RP2006043	F-NT2RP2006043	0. 50
	NT2RP2006052		0. 74
	NT2RP2006069		0. 85
40	NT2RP2006071	F-NT2RP2006071	0.94
	NT2RP2006100	F-NT2RP2006100	0. 42
	NT2RP2006106	F-NT2RP2006106	0.94
	NT2RP2006141	F-NT2RP2006141	0.92
45	NT2RP2006186	F-NT2RP2006186	0.94
	NT2RP2006196	F-NT2RP2006196	0. 32
	NT2RP2006200	F-NT2RP2006200	0.70
	NT2RP2006219	F-NT2RP2006219	0.94
50	NT2RP2006237	F-NT2RP2006237	0. 59
	NT2RP2006238	F-NT2RP2006238	0. 47
	NT2RP2006275	F-NT2RP2006275	0.89
	NT2RP2006312	F-NT2RP2006312	0. 75
55	NT2RP2006333	F-NT2RP2006333	0.73
	NT2RP2006334	F-NT2RP2006334	0. 72

	NT2RP2006365 F-NT2RP2006365	0. 48
	NT2RP2006436 F-NT2RP2006436	0. 75
	NT2RP2006441 F-NT2RP2006441	0.61
5	NT2RP2006456 F-NT2RP2006456	0. 60
	NT2RP2006464 F-NT2RP2006464	0. 51
	NT2RP2006472 F-NT2RP2006472	0. 76
	NT2RP2006565 F-NT2RP2006565	0. 34
10	NT2RP2006571 F-NT2RP2006571	0. 94
	NT2RP2006573 F-NT2RP2006573	0. 35
	NT2RP3000002 FNT2RP3000002	0.40
	NT2RP3000031 F-NT2RP3000031	0. 49
15	NT2RP3000046 F-NT2RP3000046	0.90
	NT2RP3000047 F-NT2RP3000047	0. 69
	NT2RP3000050 F-NT2RP3000050	
	NT2RP3000068 F-NT2RP3000068	0. 94
20	NT2RP3000072 F-NT2RP3000072	0. 34
	NT2RP3000080 F-NT2RP3000080	0. 32
	NT2RP3000085 F-NT2RP3000085	0. 92
	NT2RP3000092 F-NT2RP3000092	0.40
25	NT2RP3000134 F-NT2RP3000134	0.42
	NT2RP3000142 F-NT2RP3000142	0. 58
	NT2RP3000149 F-NT2RP3000149	0.83
	NT2RP3000197 F-NT2RP3000197	0.59
30	NT2RP3000207 F-NT2RP3000207	0. 48
50	NT2RP3000220 F-NT2RP3000220	0.75
	NT2RP3000251 F-NT2RP3000251	0.82
	NT2RP3000252 F-NT2RP3000252	0.88
25	NT2RP3000255 F-NT2RP3000255	0.94
35	NT2RP3000267 F-NT2RP3000267	0.92
	NT2RP3000299 F-NT2RP3000299	0.94
	NT2RP3000312 F-NT2RP3000312	0. 57
	NT2RP3000320 F-NT2RP3000320	0. 94
40	NT2RP3000324 F-NT2RP3000324	0. 32
	NT2RP3000333 F-NT2RP3000333	0. 94
	NT2RP3000348 F-NT2RP3000348	0.32
	NT2RP3000350 F-NT2RP3000350	0. 67
45	NT2RP3000359 F-NT2RP3000359	0. 94
	NT2RP3000361 F-NT2RP3000361	0.94
	NT2RP3000366 F-NT2RP3000366	0. 94
	NT2RP3000393 F-NT2RP3000393	0. 53
50	NT2RP3000397 F-NT2RP3000397	0.89
	NT2RP3000403 F-NT2RP3000403	0.87
	NT2RP3000439 F-NT2RP3000439	0. 77
	NT2RP3000456 F-NT2RP3000456	0. 88
55	NT2RP3000484 F-NT2RP3000484	0. 59
	NT2RP3000487 F-NT2RP3000487	0. 32

	NT2RP3000526	F-NT2RP3000526	0.32
	NT2RP3000527	F-NT2RP3000527	0.69
	NT2RP3000531	F-NT2RP3000531	0.56
5	NT2RP3000562	F-NT2RP3000562	0.44
	NT2RP3000590	F-NT2RP3000590	0. 88
	NT2RP3000592	F-NT2RP3000592	0. 92
	NT2RP3000596	F-NT2RP3000596	0.50
10	NT2RP3000599	F-NT2RP3000599	0. 69
	NT2RP3000603	F-NT2RP3000603	0.94
	NT2RP3000605	F-NT2RP3000605	0.92
	NT2RP3000624	F-NT2RP3000624	0.71
15	NT2RP3000632	F-NT2RP3000632	0.43
	NT2RP3000644	F-NT2RP3000644	0.94
	NT2RP3000661	F-NT2RP3000661,	0.60
	NT2RP3000665	F-NT2RP3000665	0.46
20	NT2RP3000690	F-NT2RP3000690	0.79
	NT2RP3000739	F-NT2RP3000739	0.94
	NT2RP3000753	F-NT2RP3000753	0.42
	NT2RP3000759	F-NT2RP3000759	0. 94
25	NT2RP3000825	F-NT2RP3000825	0.73
	NT2RP3000826	F-NT2RP3000826	0.94
	NT2RP3000836	F-NT2RP3000836	0. 39
	NT2RP3000841	F-NT2RP3000841	0. 55
30	NT2RP3000845	F-NT2RP3000845	0.38
	NT2RP3000847	F-NT2RP3000847	0. 67
	NT2RP3000850	F-NT2RP3000850	0. 59
	NT2RP3000852	F-NT2RP3000852	0. 39
35	NT2RP3000859	F-NT2RP3000859	0.83
	NT2RP3000868	F-NT2RP3000868	0.49
	NT2RP3000869	F-NT2RP3000869	0. 94
	NT2RP3000901	F-NT2RP3000901	0.34
40	NT2RP3000917	F-NT2RP3000917	0. 73
40	NT2RP3000919	F-NT2RP3000919	0. 94
	NT2RP3000968	F-NT2RP3000968	0. 94
	NT2RP3000980	F-NT2RP3000980	0.61
	NT2RP3000994 NT2RP3001004	F-NT2RP3000994	0.45
45	NT2RP3001004	F-NT2RP3001004	0.86
	NT2RP3001055	F-NT2RP3001055	0.38
	NT2RP3001081	F-NT2RP3001057 F-NT2RP3001081	0.84
	NT2RP3001084	F-NT2RP3001084	0. 57 0. 94
50	NT2RP3001084	F-NT2RP3001084	0. 94
	NT2RP3001098	F-NT2RP3001096	0. 94
	NT2RP3001107	F-NT2RP3001107	0. 67
	NT2RP3001111	F-NT2RP3001111	0. 87
55	NT2RP3001111	F-NT2RP3001113	0. 40
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	NT2RP3001116	E-NTORROOM 110	0 70
		F-NT2RP3001116	0. 73
	NT2RP3001119	F-NT2RP3001119	0.62
5	NT2RP3001120	F-NT2RP3001120	0. 59
	NT2RP3001126	F-NT2RP3001126	
	NT2RP3001133	F-NT2RP3001133	0.89
	NT2RP3001140	F-NT2RP3001140	0. 59
	NT2RP3001147	F-NT2RP3001147	0.94
10	NT2RP3001150	F-NT2RP3001150	0. 92
	NT2RP3001155	F-NT2RP3001155	0.94
	NT2RP3001176	F-NT2RP3001176	0.82
	NT2RP3001214	F-NT2RP3001214	0.41
15	NT2RP3001216	F-NT2RP3001216	0.32
	NT2RP3001221	F-NT2RP3001221	0.46
	NT2RP3001236	F-NT2RP3001236,	0.79
	NT2RP3001239	F-NT2RP3001239	0.94
20	NT2RP3001253	F-NT2RP3001253	0.59
	NT2RP3001260	F-NT2RP3001260	0.94
	NT2RP3001268	F-NT2RP3001268	0.75
	NT2RP3001272	F-NT2RP3001272	0.94
25	NT2RP3001274	F-NT2RP3001274	0.94
	NT2RP3001297	F-NT2RP3001297	0.94
	NT2RP3001307	F-NT2RP3001307	0.87
	NT2RP3001325	F-NT2RP3001325	0. 71
	NT2RP3001338	F-NT2RP3001338	0.90
30	NT2RP3001355	F-NT2RP3001355	0.40
	NT2RP3001356	F-NT2RP3001356	0. 88
	NT2RP3001384	F-NT2RP3001384	0.91
	NT2RP3001392	F-NT2RP3001392	0.42
35	NT2RP3001396	F-NT2RP3001396	
	NT2RP3001398	F-NT2RP3001398	0. 94
	NT2RP3001399	F-NT2RP3001399	0.94
	NT2RP3001407	F-NT2RP3001407	0. 54
40	NT2RP3001420	F-NT2RP3001420	0.74
	NT2RP3001426	F-NT2RP3001426	0. 92
	NT2RP3001427		
		F-NT2RP3001428	0. 94
45	NT2RP3001447	F-NT2RP3001447	0. 57
	NT2RP3001449	F-NT2RP3001449	0. 62
	NT2RP3001453	F-NT2RP3001453	0. 94
	NT2RP3001457	F-NT2RP3001457	0.89
50	NT2RP3001472	F-NT2RP3001472	0.82
50	NT2RP3001490	F-NT2RP3001490	0. 53
	NT2RP3001495	F-NT2RP3001495	0. 94
	NT2RP3001497	F-NT2RP3001497	0. 40
	NT2RP3001529	F-NT2RP3001497	0. 94
55	NT2RP3001538	F-NT2RP3001538	0. 94
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	NT-2RP3001554	F-NT2RP3001554	0.91
	NT2RP3001587	F-NT2RP3001587	0.94
	NT2RP3001608	F-NT2RP3001608	0.94
5	NT2RP3001621	F-NT2RP3001621	0. 78
	NT2RP3001629	F-NT2RP3001629	0. 42
	NT2RP3001642	F-NT2RP3001642	0.94
	NT2RP3001646	F-NT2RP3001646	0. 94
10	NT2RP3001671	F-NT2RP3001671	0. 77
	NT2RP3001672	F-NT2RP3001672	0.88
	NT2RP3001676	F-NT2RP3001676	0.88
	NT2RP3001678	F-NT2RP3001678	0. 44
15	NT2RP3001679	F-NT2RP3001679	0. 44
	NT2RP3001688	F-NT2RP3001688	0. 67
	NT2RP3001690	F-NT2RP3001690,	
	NT2RP3001698	F-NT2RP3001698	0. 86
20	NT2RP3001708	F-NT2RP3001708	0. 38
20	NT2RP3001712	F-NT2RP3001712	0. 35
	NT2RP3001716	F-NT2RP3001716	0. 73
	NT2RP3001724		0. 77
	NT2RP3001727		0. 84
25	NT2RP3001730	F-NT2RP3001730	0.77
	NT2RP3001739	F-NT2RP3001739	0. 78
	NT2RP3001753	F-NT2RP3001753	0. 79
	NT2RP3001764	F-NT2RP3001764	0. 73
30	NT2RP3001777	F-NT2RP3001777	0.87
	NT2RP3001792	F-NT2RP3001792	0.86
	NT2RP3001799	F-NT2RP3001799	0. 79
	NT2RP3001819	F-NT2RP3001819	0. 73
35	NT2RP3001844	F-NT2RP3001844	0. 34
	NT2RP3001854	F-NT2RP3001854	0. 94
	NT2RP3001855	F-NT2RP3001855	0. 88
	NT2RP3001857	F-NT2RP3001857	0. 41
40	NT2RP3001896	F-NT2RP3001896	0. 94
	NT2RP3001915	F-NT2RP3001915	0. 42
	NT2RP3001929	F-NT2RP3001929	0. 49
	NT2RP3001931	F-NT2RP3001931	0. 94
45	NT2RP3001938	F-NT2RP3001938	0. 32
45	NT2RP3001943	F-NT2RP3001943	0. 70
	NT2RP3001944	F-NT2RP3001944	0. 94
	NT2RP3001969	F-NT2RP3001969	0. 54
-0	NT2RP3002007	F-NT2RP3002007	0. 56
50	NT2RP3002014	F-NT2RP3002014	0. 94
	NT2RP3002033	F-NT2RP3002033	0. 94
	NT2RP3002035	F-NT2RP3002045	0. 94
	NT2RP3002054	F-NT2RP3002054	0. 91
55	NT2RP3002062	F-NT2RP3002062	0. 62
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	NT2RP3002063	E-NTODDOOOGO	0.24
	NT2RP3002081	F-NT2RP3002063	0.34
		F-NT2RP3002081	0. 57
5	NT2RP3002097	F-NT2RP3002097	0. 72
•	NT2RP3002102	F-NT2RP3002102	0.90
	NT2RP3002108	F-NT2RP3002108	0.74
	NT2RP3002142	F-NT2RP3002142	0. 61
	NT2RP3002146	F-NT2RP3002146	0. 91
10	NT2RP3002147	F-NT2RP3002147	0. 37
	NT2RP3002151	F-NT2RP3002151	0. 90
	NT2RP3002163	F-NT2RP3002163	0. 76
	NT2RP3002165	F-NT2RP3002165	0.89
15	NT2RP3002181	F-NT2RP3002181	0.78
	NT2RP3002244	F-NT2RP3002244	0.81
	NT2RP3002248	F-NT2RP3002248,	0.36
	NT2RP3002255	F-NT2RP3002255	0.53
20	NT2RP3002273	F-NT2RP3002273	0.90
	NT2RP3002276	F-NT2RP3002276	0.39
	NT2RP3002303	F-NT2RP3002303	0.40
	NT2RP3002304	F-NT2RP3002304	0.44
25	NT2RP3002330	F-NT2RP3002330	0.94
20	NT2RP3002343	F-NT2RP3002343	0.94
	NT2RP3002351	F-NT2RP3002351	0.94
	NT2RP3002377	F-NT2RP3002377	0. 52
	NT2RP3002399	F-NT2RP3002399	0. 91
30	NT2RP3002402	F-NT2RP3002402	0.33
	NT2RP3002455	F-NT2RP3002455	0.42
	NT2RP3002484	F-NT2RP3002484	0. 52
	NT2RP3002501	F-NT2RP3002501	0. 57
35	NT2RP3002512	F-NT2RP3002512	0. 63
	NT2RP3002529	F-NT2RP3002529	0.74
	NT2RP3002545	F-NT2RP3002545	0. 94
	NT2RP3002549	F-NT2RP3002549	0, 94
40	NT2RP3002566	F-NT2RP3002566	0.74
	NT2RP3002587	F-NT2RP3002587	0. 94
	NT2RP3002590	F-NT2RP3002590	
	NT2RP3002602	F-NT2RP3002602	0.39
45	NT2RP3002603	F-NT2RP3002603	0. 46
	NT2RP3002628	F-NT2RP3002628	0. 31
	NT2RP3002631	F-NT2RP3002631	0. 79
	NT2RP3002650	F-NT2RP3002650	0. 94
	NT2RP3002659	F-NT2RP3002659	0. 93
50	NT2RP3002660	F-NT2RP3002660	0. 46
	NT2RP3002663	F-NT2RP3002663	0. 79
	NT2RP3002671	F-NT2RP3002671	0.73
	NT2RP3002682	F-NT2RP3002682	0. 52
55	NT2RP3002688	F-NT2RP3002688	0. 82
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	NT2RP3002701	F-NT2RP3002701	0.62
	NT2RP3002763	F-NT2RP3002763	0. 83
	NT2RP3002785	F-NT2RP3002785	0. 33
5	NT2RP3002818	F-NT2RP3002818	0. 91
	NT2RP3002861	F-NT2RP3002861	0. 31
	NT2RP3002869	F-NT2RP3002869	0.47
	NT2RP3002876	F-NT2RP3002876	0. 58
10	NT2RP3002877	F-NT2RP3002877	0. 41
	NT2RP3002909	F-NT2RP3002909	0. 52
	NT2RP3002911	F-NT2RP3002911	0. 73
	NT2RP3002948	F-NT2RP3002948	0. 60
15	NT2RP3002953	F-NT2RP3002953	0.37
	NT2RP3002969	F-NT2RP3002969	0. 47
	NT2RP3002972	F-NT2RP3002972,	0. 87
	NT2RP3002985	F-NT2RP3002985	0. 55
20	NT2RP3002988	F-NT2RP3002988	0.82
	NT2RP3003008	F-NT2RP3003008	0.94
	NT2RP3003061	F-NT2RP3003061	0.93
	NT2RP3003068	F-NT2RP3003068	0. 45
25	NT2RP3003071	F-NT2RP3003071	0.76
	NT2RP3003078	F-NT2RP3003078	0.40
	NT2RP3003101	F-NT2RP3003101	0. 41
	NT2RP3003139	F-NT2RP3003139	0.43
30	NT2RP3003145	F-NT2RP3003145	0. 94
	NT2RP3003150	F-NT2RP3003150	0. 94
	NT2RP3003157	F-NT2RP3003157	0. 81
	NT2RP3003185	F-NT2RP3003185	0. 87
35	NT2RP3003193	F-NT2RP3003193	0. 94
	NT2RP3003197	F-NT2RP3003197	0.74
	NT2RP3003203	F-NT2RP3003203	0. 94
	NT2RP3003204	F-NT2RP3003204	0. 74
40	NT2RP3003210	F-NT2RP3003210	0.45
40	NT2RP3003212 NT2RP3003230	F-NT2RP3003212 F-NT2RP3003230	0.51
	NT2RP3003242	F-NT2RP3003242	0. 94 0. 92
	NT2RP3003251	F-NT2RP3003251	0. 92
_	NT2RP3003278	F-NT2RP3003278	0. 83
45	NT2RP3003282	F-NT2RP3003282	0. 67
	NT2RP3003290	F-NT2RP3003290	0. 62
	NT2RP3003301	F-NT2RP3003301	0. 49
	NT2RP3003302	F-NT2RP3003302	0. 32
50	NT2RP3003311	F-NT2RP3003311	0. 34
	NT2RP3003313	F-NT2RP3003313	0. 88
	NT2RP3003327	F-NT2RP3003327	0. 88
	NT2RP3003330	F-NT2RP3003330	0. 94
55	NT2RP3003344	F-NT2RP3003344	0. 67
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	NT2RP3003353	F-NT2RP3003353	0. 65
	NT2RP3003377	F-NT2RP3003377	0. 87
	NT2RP3003385	F-NT2RP3003385	0. 94
5	NT2RP3003409	F-NT2RP3003409	0.41
	NT2RP3003411	F-NT2RP3003411	0, 47
	NT2RP3003427	F-NT2RP3003427	0. 33
	NT2RP3003490	F-NT2RP3003490	0. 56
10	NT2RP3003491	F-NT2RP3003491	0. 37
	NT2RP3003543	F-NT2RP3003543	0. 92
	NT2RP3003552	F-NT2RP3003552	0. 64
	NT2RP3003555	F-NT2RP3003555	0. 94
15	NT2RP3003564	F-NT2RP3003564	0.86
	NT2RP3003589	F-NT2RP3003589	0.88
	NT2RP3003621	F-NT2RP3003621,	0.49
	NT2RP3003656	F-NT2RP3003656	0. 77
20	NT2RP3003659	F-NT2RP3003659	0.32
	NT2RP3003672	F-NT2RP3003672	0.49
	NT2RP3003680	F-NT2RP3003680	0.94
	NT2RP3003686	F-NT2RP3003686	0.78
25	NT2RP3003701	F-NT2RP3003701	0.36
	NT2RP3003716	F-NT2RP3003716	0.34
	NT2RP3003726	F-NT2RP3003726	0.72
	NT2RP3003795	F-NT2RP3003795	0.34
30	NT2RP3003805	F-NT2RP3003805	0. 74
30	NT2RP3003809	F-NT2RP3003809	0. 73
	NT2RP3003825	F-NT2RP3003825	0.94
	NT2RP3003831	F-NT2RP3003831	0. 52
	NT2RP3003833	F-NT2RP3003833	0.31
35	NT2RP3003846	F-NT2RP3003846	0.54
	NT2RP3003870	F-NT2RP3003870	0. 91
	NT2RP3003876	F-NT2RP3003876	0. 42
	NT2RP3003914	F-NT2RP3003914	0.94
40	NT2RP3003918	F-NT2RP3003918	0. 90
	NT2RP3003932	F-NT2RP3003932	0.64
	NT2RP3003992	F-NT2RP3003992	0. 94
	NT2RP3004013 NT2RP3004041	F-NT2RP3004013	0.60
45	NT2RP3004041	F-NT2RP3004041 F-NT2RP3004051	0. 52 0. 75
	NT2RP3004051	F-NT2RP3004078	0. 73
	NT2RP3004078	F-NT2RP3004093	0. 73
	NT2RP3004093	F-NT2RP3004093	0. 94
50	NT2RP3004125	F-NT2RP3004155	0. 89
	NT2RP3004189	F-NT2RP3004189	0. 46
	NT2RP3004206	F-NT2RP3004206	0. 59
	NT2RP3004207	F-NT2RP3004207	0. 39
55	NT2RP3004209	F-NT2RP3004209	0. 54
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	NT2RP3004242	F-NT2RP3004242	0.61
	NT2RP3004246	F-NT2RP3004246	0.91
	NT2RP3004253	F-NT2RP3004253	0.41
5	NT2RP3004258	F-NT2RP3004258	0.41
	NT2RP3004262	F-NT2RP3004262	0. 56
	NT2RP3004282	F-NT2RP3004282	0.90
	NT2RP3004332	F-NT2RP3004332	0. 53
10	NT2RP3004341	F-NT2RP3004341	0.40
	NT2RP3004348	F-NT2RP3004348	0.94
	NT2RP3004378	F-NT2RP3004378	0.40
	NT2RP3004424	F-NT2RP3004424	0. 75
15	NT2RP3004428	F-NT2RP3004428	0, 85
	NT2RP3004451	F-NT2RP3004451	0. 85
	NT2RP3004454	F-NT2RP3004454,	
	NT2RP3004466	F-NT2RP3004466	0.80
20	NT2RP3004472	F-NT2RP3004472	0.46
20	NT2RP3004480	F-NT2RP3004480	0. 77
	NT2RP3004490	F-NT2RP3004490	0. 65
·	NT2RP3004498	F-NT2RP3004498	0.91
25	NT2RP3004504	F-NT2RP3004504	0.88
25	NT2RP3004507	F-NT2RP3004507	0.86
	NT2RP3004534	F-NT2RP3004534	0.32
	NT2RP3004539	F-NT2RP3004539	0.76
	NT2RP3004544	F-NT2RP3004544	0.94
30	NT2RP3004566	F-NT2RP3004566	0.94
	NT2RP3004569	F-NT2RP3004569	0.69
	NT2RP3004572	F-NT2RP3004572	0. 65
	NT2RP3004578	F-NT2RP3004578	0, 52
35	NT2RP3004594	F-NT2RP3004594	0. 39
	NT2RP3004617	F-NT2RP3004617	0. 94
	NT2RP3004618	F-NT2RP3004618	0.41
	NT2RP3004669	F-NT2RP3004669	0. 82
40	NT2RP3004670	F-NT2RP3004670	0. 32
	NT2RP4000008	F-NT2RP4000008	0.94
	NT2RP4000051	F-NT2RP4000051	0.94
	NT2RP4000109	F-NT2RP4000109	0.84
45	NT2RP4000111	F-NT2RP4000111	0. 49
	NT2RP4000129	F-NT2RP4000129	0.94
	NT2RP4000147	F-NT2RP4000147	0. 92
	NT2RP4000150	F-NT2RP4000150	0. 41
50	NT2RP4000151	F-NT2RP4000151	0. 33
	NT2RP4000159	F-NT2RP4000159	0. 75
	NT2RP4000185	F-NT2RP4000185	0. 94
	NT2RP4000210	F-NT2RP4000210	0. 83
55	NT2RP4000212	F-NT2RP4000212	0.94
	NT2RP4000243	F-NT2RP4000243	0. 94

	NT2RP4000246	F-NT2RP4000246	0. 59
	NT2RP4000259	F-NT2RP4000259	0. 94
		F-NT2RP4000263	0. 62
5	NT2RP4000290	F-NT2RP4000290	0. 92
	NT2RP4000312		0. 39
	NT2RP4000323	F-NT2RP4000323	0. 73
	NT2RP4000355	F-NT2RP4000355	0. 94
10	NT2RP4000360	F-NT2RP4000360	0. 55
	NT2RP4000367		0. 60
	NT2RP4000370		0. 93
	NT2RP4000376		0. 88
15	NT2RP4000381	F-NT2RP4000381	0. 39
	NT2RP4000398		0. 64
	NT2RP4000415		
	NT2RP4000417	•	0. 33
20	NT2RP4000448		0. 33
20	NT2RP4000449		0.33
	NT2RP4000455		0. 37
	NT2RP4000457		0. 36
	NT2RP4000480	F-NT2RP4000480	0. 88
25	NT2RP4000481	F-NT2RP4000481	0. 51
	NT2RP4000498	F-NT2RP4000498	0. 64
	NT2RP4000500	F-NT2RP4000500	0. 67
	NT2RP4000518	F-NT2RP4000518	0. 34
30	NT2RP4000524	F-NT2RP4000524	0. 94
	NT2RP4000528	F-NT2RP4000528	0. 65
	NT2RP4000541	F-NT2RP4000541	0. 44
	NT2RP4000556	F-NT2RP4000556	0. 69
35	NT2RP4000560	F-NT2RP4000560	0. 67
	NT2RP4000588	F-NT2RP4000588	0. 94
	NT2RP4000614	F-NT2RP4000614	0. 94
	NT2RP4000638	F-NT2RP4000638	0. 74
40	NT2RP4000648	F-NT2RP4000648	0. 40
	NT2RP4000657	F-NT2RP4000657	0. 94
	NT2RP4000704	F-NT2RP4000704	0. 69
	NT2RP4000713	F-NT2RP4000713	0. 93
45	NT2RP4000724	F-NT2RP4000724	0. 32
40	NT2RP4000728	F-NT2RP4000728	0. 34
	NT2RP4000737	F-NT2RP4000737	0. 48
	NT2RP4000739	F-NT2RP4000739	0. 40
	NT2RP4000781	F-NT2RP4000781	0. 94
50	NT2RP4000787	F-NT2RP4000787	0. 94
	NT2RP4000817	F-NT2RP4000817	0.60
	NT2RP4000833	F-NT2RP4000833	0. 35
	NT2RP4000837	F-NT2RP4000837	0.80
55	NT2RP4000839	F-NT2RP4000839	0. 33

	NT2RP4000855	F-NT2RP4000855	0. 74
	NT2RP4000865	F-NT2RP4000865	0. 77
	NT2RP4000878	F-NT2RP4000878	0. 79
5	NT2RP4000879	F-NT2RP4000879	0. 41
	NT2RP4000907	F-NT2RP4000907	0. 44
	NT2RP4000918	F-NT2RP4000918	0. 88
	NT2RP4000925	F-NT2RP4000925	0. 89
10	NT2RP4000927	F-NT2RP4000927	0. 81
	NT2RP4000928	F-NT2RP4000928	0. 39
	NT2RP4000929	F-NT2RP4000929	0. 34
	NT2RP4000955	F-NT2RP4000955	0. 77
15	NT2RP4000973	F-NT2RP4000973	0.36
	NT2RP4000975	F-NT2RP4000975	0. 87
	NT2RP4000979	F-NT2RP4000979	0.50
	NT2RP4000984	F-NT2RP4000984	0. 66
20	NT2RP4000989	F-NT2RP4000989	0. 94
	NT2RP4000997	F-NT2RP4000997	0. 68
	NT2RP4001004	F-NT2RP4001004	0.40
	NT2RP4001006	F-NT2RP4001006	0. 91
25	NT2RP4001010	F-NT2RP4001010	0. 64
25	NT2RP4001029	F-NT2RP4001029	0.77
	NT2RP4001041	F-NT2RP4001041	0. 79
	NT2RP4001057	F-NT2RP4001057	0. 65
	NT2RP4001064	F-NT2RP4001064	0. 94
30	NT2RP4001078	F-NT2RP4001078	0. 39
	NT2RP4001079	F-NT2RP4001079	0. 94
	NT2RP4001080	F-NT2RP4001080	0. 58
	NT2RP4001086	F-NT2RP4001086	0. 78
35	NT2RP4001095	F-NT2RP4001095	0. 80
	NT2RP4001100	F-NT2RP4001100	0. 79
	NT2RP4001117	F-NT2RP4001117	0. 60
	NT2RP4001122	F-NT2RP4001122	0. 82
40	NT2RP4001126	F-NT2RP4001126	0. 55
	NT2RP4001138	F-NT2RP4001138	0. 94
	NT2RP4001143		0. 41
	NT2RP4001148	F-NT2RP4001148	0. 37
45	NT2RP4001149		0. 50
	NT2RP4001150	F-NT2RP4001150	0. 94
	NT2RP4001159	F-NT2RP4001159	0. 78
	NT2RP4001174	F-NT2RP4001174	0. 94
50	NT2RP4001206	F-NT2RP4001206	0.80
	NT2RP4001207	F-NT2RP4001207	0. 38
	NT2RP4001210	F-NT2RP4001210	0. 94
	NT2RP4001213	F-NT2RP4001213	0. 46
55	NT2RP4001219	F-NT2RP4001219	0. 38
	NT2RP4001228	F-NT2RP4001228	0. 61

	NT2RP4001235	F-NT2RP4001235	0. 88
	NT2RP4001256	F-NT2RP4001256	0. 60
	NT2RP4001260	F-NT2RP4001260	0. 94
5	NT2RP4001274	F-NT2RP4001274	0. 82
	NT2RP4001276	F-NT2RP4001276	0. 68
	NT2RP4001313	F-NT2RP4001313	0. 77
	NT2RP4001315	F-NT2RP4001315	0. 87
10	NT2RP4001336	F-NT2RP4001336	0. 48
	NT2RP4001339	F-NT2RP4001339	0. 41
	NT2RP4001343	F-NT2RP4001343	0. 63
	NT2RP4001345	F-NT2RP4001345	0. 89
15	NT2RP4001351	F-NT2RP4001351	0. 87
	NT2RP4001353	F-NT2RP4001353	0. 72
	NT2RP4001372	F-NT2RP4001372,	0. 78
	NT2RP4001373	F-NT2RP4001373	0.60
20	NT2RP4001375	F-NT2RP4001375	0.43
	NT2RP4001379	F-NT2RP4001379	0.77
	NT2RP4001389	F-NT2RP4001389	0. 41
	NT2RP4001414	F-NT2RP4001414	0.90
25	NT2RP4001433	F-NT2RP4001433	0.39
	NT2RP4001442	F-NT2RP4001442	0.59
	NT2RP4001447	F-NT2RP4001447	0. 33
	NT2RP4001474	F-NT2RP4001474	0.89
30	NT2RP4001483	F-NT2RP4001483	0. 94
30	NT2RP4001498	F-NT2RP4001498	0. 59
	NT2RP4001502	F-NT2RP4001502	0.61
	NT2RP4001507	F-NT2RP4001507	0. 91
25	NT2RP4001524	F-NT2RP4001524	0.61
35	NT2RP4001529	F-NT2RP4001529	0.75
	NT2RP4001547	F-NT2RP4001547	0. 77
	NT2RP4001551	F-NT2RP4001551	0. 79
	NT2RP4001555	F-NT2RP4001555	0. 79
40	NT2RP4001567	F-NT2RP4001567	0. 39
	NT2RP4001568	F-NT2RP4001568	0. 45
	NT2RP4001571	F-NT2RP4001571	0. 76
	NT2RP4001574	F-NT2RP4001574	0. 91
45	NT2RP4001575	F-NT2RP4001575	0. 57
	NT2RP4001592	F-NT2RP4001592	0.71
	NT2RP4001610	F-NT2RP4001610	0.94
	NT2RP4001614	F-NT2RP4001614	0. 73
50	NT2RP4001634	F-NT2RP4001634	0.40
	NT2RP4001638	F-NT2RP4001638	0. 37
	NT2RP4001644	F-NT2RP4001644	0. 94
	NT2RP4001656	F-NT2RP4001656	0. 92
55	NT2RP4001677	F-NT2RP4001677	0. 47
	NT2RP4001679	F-NT2RP4001679	0. 54

	NT2RP4001696	F-NT2RP4001696	0. 62
	NT2RP4001725	F-NT2RP4001725	0. 94
	NT2RP4001730	F-NT2RP4001730	0. 94
5	NT2RP4001739	F-NT2RP4001739	0. 82
	NT2RP4001753		0. 66
	NT2RP4001760		0. 39
	NT2RP4001790		0. 64
10	NT2RP4001803		0. 87
	NT2RP4001822		0. 84
	NT2RP4001823		0. 56
	NT2RP4001828		0. 45
15	NT2RP4001838		0. 43
•	NT2RP4001841		0. 58
	NT2RP4001849	F-NT2RP4001849,	
	NT2RP4001861	•	0.80
00	NT2RP4001893		0. 94
20	NT2RP4001896		0. 94
	NT2RP4001990		0. <del>9</del> 3 0. 89
	NT2RP4001907		0. <i>03</i> 0. 77
	NT2RP4001927		0. 77 0. 94
25	NT2RP4001938		0. 74
	NT2RP4001940	•	
	NT2RP4001953		0. 45 0. 49
	NT2RP4001966		
30	NT2RP4001975		0. 86 0. 94
	NT2RP4001975		
	NT2RP4002018		0. 67
	NT2RP4002047		0. 67
35	NT2RP4002058		0. 42
	NT2RP4002038		0. 78 0. 94
	NT2RP4002071		0. <del>34</del> 0. 38
	NT2RP4002078		0. 55 0. 55
40	NT2RP4002081		0. 86
10	NT2RP4002408		0. 64
	NT2RP4002791		0. 84
	NT2RP4002888		
	NT2RP5003459		0. 31 0. 94
45	NT2RP5003461		
	NT2RP5003461		0. 94 0. 44
	NT2RP5003492		0. 57
50	NT2RP5003500		0. 94
	NT2RP5003506		0. 94
	NT2RP5003512		0. 42
	NT2RP5003522		0. 94
55	NT2RP5003524		0. 58
	NT2RP5003534	F-NT2RP5003534	0. 32

	OVARC1000001	F-0VARC1000001	0.74
	OVARC1000004	F-0VARC1000004	0. 90
	0VARC1000006	F-0VARC1000006	0. 94
5	OVARC1000013	F-0VARC1000013	0.48
	0VARC1000014	F-0VARC1000014	0. 69
	0VARC1000035	F-0VARC1000035	0. 59
	0VARC1000060	F-0VARC1000060	0.76
10	OVARC1000087	F-0VARC1000087	0. 31
	0VARC1000091	F-0VARC1000091	0. 88
	0VARC1000106	F-0VARC1000106	0.88
	0VARC1000109	F-0VARC1000109	0.94
15	0VARC1000113	F-0VARC1000113	0.40
	0VARC1000114	F-0VARC1000114	0. 53
	0VARC1000139	F-0VARC1000139,	0.88
	0VARC1000148	F-0VARC1000148	0.94
20	0VARC1000151	F-0VARC1000151	0.34
	OVARC1000168	F-0VARC1000168	0.56
	0VARC1000209	F-0VARC1000209	0.94
	OVARC1000212	F-0VARC1000212	0. 87
25	OVARC1000241	F-0VARC1000241	0.59
25	0VARC1000288	F-0VARC1000288	0.87
	0VARC1000304	F-0VARC1000304	0.94
	OVARC1000309	F-0VARC1000309	0.76
	OVARC1000321	F-0VARC1000321	0.90
30	0VARC1000326	F-0VARC1000326	0.46
	0VARC1000335	F-0VARC1000335	0.55
	0VARC1000347	F-0VARC1000347	0.43
	0VARC1000384	F-0VARC1000384	0.90
35	0VARC1000408	F-0VARC1000408	0.42
	OVARC1000411	F-0VARC1000411	0.65
	0VARC1000420	F-0VARC1000420	0.84
	0VARC1000437	F-0VARC1000437	0. 94
40	0VARG1000440	F-0VARC1000440	0. 94
	0VARC1000442	F-0VARC1000442	0.94
	0VARC1000443	F-0VARC1000443	
	0VARC1000461	F-0VARC1000461	0. 89
45	OVARC1000465	F-0VARC1000465	0. 75
	0VARC1000466	F-0VARC1000466	0.94
	0VARC1000473	F-0VARC1000473	0.31
	0VARC1000479	F-0VARC1000479	0. 42
50	OVARC1000520	F-0VARC1000520	0.94
	0VARC1000556	F-0VARC1000556	0. 94
	OVARC1000557	F-0VARC1000557	0. 72
	OVARC1000564	F-0VARC1000564	0. 94
55	0VARC1000576	F-0VARC1000576	0. 40
	OVARC1000588	F-0VARC1000588	0. 45

	0VARC1000605	F-0VARC1000605	0. 58
	0VARC1000622	F-0VARC1000622	0.80
	0VARC1000640	F-0VARC1000640	0. 35
5	OVARC1000649	F-0VARC1000649	
	0VARC1000661	F-0VARC1000661	0.87
	0VARC1000681	F-0VARC1000681	0.84
	0VARC1000682	F-0VARC1000682	0. 38
10	0VARC1000689	F-0VARC1000689	0.36
	0VARC1000003	F-0VARC1000703	0. 92
	0VARC1000703	F-0VARC1000722	0. 94
	OVARC1000722	F-0VARC1000722	0. 90
15	0VARC1000736	F-0VARC1000746	0. 42
13	0VARC1000748	F-0VARC1000746	0. 42
	OVARC1000781 OVARC1000787	F-0VARC1000781,	
		F-0VARC1000787	0, 41
20	0VARC1000800	F-0VARC1000800	0. 85
	0VARC1000834	F-0VARC1000834	0. 62
	OVARC1000846	F-0VARC1000846	0.49
	0VARC1000850	F-0VARC1000850	0. 94
25	OVARC1000862	F-0VARC1000862	0. 54
	OVARC1000876	F-0VARC1000876	0. 74
	OVARC1000883	F-0VARC1000883	0. 55
	OVARC1000885	F-0VARC1000885	0.94
30	OVARC1000886	F-0VARC1000886	0.61
	OVARC1000890	F-0VARC1000890	0. 45
	0VARC1000912	F-0VARC1000912	0. 63
	OVARC1000915	F-0VARC1000915	0. 31
35	OVARC1000924	F-0VARC1000924	0. 82
35	OVARC1000945	F-0VARC1000945	0. 74
	0VARC1000959	F-0VARC1000959	0. 33
	OVARC1000964	F-0VARC1000964	0. 60
	0VARC1000984	F-0VARC1000984	0. 59
40	0VARC1000996	F-0VARC1000996	0. 83
	OVARC1000999	F-0VARC1000999	0. 94
	0VARC1001004	F-0VARC1001004	
	0VARC1001010	F-0VARC1001010	0.51
45	0VARC1001032	F-0VARC1001032	0. 32
	0VARC1001034	F-0VARC1001034	0. 78
	0VARC1001038	F-0VARC1001038	0. 93
	0VARC1001040	F-0VARC1001040	0. 68
50	0VARC1001044	F-0VARC1001044	0. 66
	0VARC1001055	F-0VARC1001055	0. 76
	OVARC1001065	F-0VARC1001065	0. 31
	0VARC1001068	F-0VARC1001068	0. 63
55	0VARC1001074	F-0VARC1001074	0. 82
	0VARC1001092	F-0VARC1001092	0. 50

	OVARC1001107	F-0VARC1001107	0. 65
	OVARC1001113	F-0VARC1001113	0.86
	OVARC1001154	F-0VARC1001154	0.43
5	OVARC1001161	F-0VARC1001161	0, 75
	0VARC1001162	F-0VARC1001162	0.61
	OVARC1001167	F-0VARC1001167	0.31
	OVARC1001170	F-0VARC1001170	0.46
10	OVARC1001171	F-0VARC1001171	0.78
	OVARC1001173	F-0VARC1001173	0.71
	OVARC1001176	F-0VARC1001176	0.83
	OVARC1001180	F-0VARC1001180	0. 53
15	OVARC1001188	F-0VARC1001188	0. 42
	0VARC1001200	F-0VARC1001200	0. 94
	OVARC1001232	F-0VARC1001232,	
	OVARC1001243	F-0VARC1001243	0. 59
	OVARC1001244	F-0VARC1001244	0. 79
20	OVARC1001270	F-0VARC1001270	0. 48
	0VARC1001271	F-0VARC1001271	0. 67
	OVARC1001296	F-0VARC1001296	0.60
	OVARC1001306	F-0VARC1001306	0.44
25	0VARC1001329	F-0VARC1001329	0. 91
	0VARC1001341	F-0VARC1001341	0.82
	OVARC1001344	F-0VARC1001344	0. 94
	OVARC1001360	F-0VARC1001360	0. 94
30	OVARC1001369	F-0VARC1001369	0. 36
	OVARC1001372	F-0VARC1001372	0.63
	OVARC1001376	F-0VARC1001376	0. 81
	OVARC1001381	F-0VARC1001381	0. 78
35	OVARC1001391	F-0VARC1001391	0. 33
	0VARC1001399	F-0VARC1001399	0. 67
	0VARC1001417	F-0VARC1001417	0.94
	OVARC1001419	F-0VARC1001419	0.34
40	OVARC1001425	F-0VARC1001425	0. 73
	0VARC1001436	F-0VARC1001436	0. 59
	0VARC1001453	F-0VARC1001453	0.45
	OVARC1001476	F-0VARC1001476	0.85
45	OVARC1001489	F-0VARC1001489	0.35
10	OVARC1001496	F-0VARC1001496	0.90
	OVARC1001506	F-0VARC1001506	0. 59
	OVARC1001525	F-0VARC1001525	0.87
	OVARC1001555	F-0VARC1001555	0. 67
50	OVARC1001577	F-0VARC1001577	0.94
	OVARC1001610	F-0VARC1001610	0.38
	OVARC1001611	F-0VARC1001611	0.34
	OVARC1001702	F-0VARC1001702	0.88
55	OVARC1001703	F-0VARC1001703	0. 76

	OVARC1001711	F-0VARC1001711	0.82
	OVARC1001713	F-0VARC1001713	0. 82
	OVARC1001731	F-0VARC1001731	0.94
5	0VARC1001762	F-0VARC1001762	0. 59
	OVARC1001766	F-0VARC1001766	0. 94
	OVARC1001767	F-0VARC1001767	0. 70
	OVARC1001768	F-0VARC1001768	0. 32
10	OVARC1001791	F-0VARC1001791	0.74
	0VARC1001802	F-0VARC1001802	0. 76
	OVARC1001809	F-0VARC1001809	0.63
	OVARC1001820	F-0VARC1001820	0. 94
15	OVARC1001828	F-0VARC1001828	0. 78
	OVARC1001861	F-0VARC1001861	0. 94
	OVARC1001873	F-0VARC1001873,	0. 31
	OVARC1001879	F-0VARC1001879	0. 94
20	OVARC1001880	F-0VARC1001880	0, 46
20	OVARC1001916	F-0VARC1001916	0. 82
	OVARC1001942	F-0VARC1001942	0.72
	OVARC1001943	F-0VARC1001943	0.49
	OVARC1001949	F-0VARC1001949	0.46
25	OVARC1001950	F-0VARC1001950	0. 94
	OVARC1001987.	F-0VARC1001987	0.91
	OVARC1002050	F-0VARC1002050	0.85
	OVARC1002082	F-0VARC1002082	0.94
30	OVARC1002107	F-0VARC1002107	0.33
	OVARC1002112	F-0VARC1002112	0.94
	OVARC1002127	F-0VARC1002127	0.80
	OVARC1002138	F-0VARC1002138	0.46
35	OVARC1002156	F-0VARC1002156	0. 43
	OVARC1002158	F-0VARC1002158	0. 94
	OVARC1002165	F-0VARC1002165	0.88
	OVARC1002182	F-0VARC1002182	0. 94
40	PLACE1000004	F-PLACE1000004	0.43
	PLACE1000005	F-PLACE1000005	0.37
	PLACE1000007	F-PLACE1000007	0.72
	PLACE1000014	F-PLACE1000014	0.38
45	PLACE1000048	F-PLACE1000048	0. 36
	PLACE1000050	F-PLACE1000050	0. 94
	PLACE1000061	F-PLACE1000061	0. 56
	PLACE1000066	F-PLACE1000066	0. 94
50	PLACE1000081	F-PLACE1000081	0.37
	PLACE1000133	F-PLACE1000133	0. 53
	PLACE1000142	F-PLACE1000142	0.94
	PLACE1000184	F-PLACE1000184	0.73
g.c	PLACE1000185	F-PLACE1000185	0.86
55	PLACE1000213	F-PLACE1000213	0.34

	PLACE1000236	F-PLACE1000236	0. 76
	PLACE1000246	F-PLACE1000246	0.91
	PLACE1000308	F-PLACE1000308	0.40
5	PLACE1000347	F-PLACE1000347	0.48
	PLACE1000374	F-PLACE1000374	0. 58
	PLACE1000380	F-PLACE1000380	0. 94
	PLACE1000383	F-PLACE1000383	0. 39
10	PLACE1000401	F-PLACE1000401	0. 94
	PLACE1000406	F-PLACE1000406	0. 94
	PLACE1000420	F-PLACE1000420	0.82
	PLACE1000425	F-PLACE1000435	0. 32
45	PLACE1000444	F-PLACE1000444	0. 41
15	PLACE1000444 PLACE1000453	F-PLACE1000453	0. 76
	PLACE1000493		
		F-PLACE1000492,	0. 94
	PLACE1000547	F-PLACE1000547	0. 94
20	PLACE1000562	F-PLACE1000562	
	PLACE1000564	F-PLACE1000564	0.45
	PLACE1000588	F-PLACE1000588	0.67
	PLACE1000596	F-PLACE1000596	0.60
25	PLACE1000610	F-PLACE1000610	0.94
	PLACE1000611	F-PLACE1000611	0. 48
	PLACE1000636	F-PLACE1000636	0. 92
	PLACE1000653	F-PLACE1000653	0. 57
30	PLACE1000656	F-PLACE1000656	0. 94
	PLACE1000706	F-PLACE1000706	0. 94
	PLACE1000712	F-PLACE1000712	0. 78
	PLACE1000716	F-PLACE1000716	0. 94
	PLACE1000748	F-PLACE1000748	0. 52
35	PLACE1000755	F-PLACE1000755	0.87
	PLACE1000769	F-PLACE1000769	0. 94
	PLACE1000785	F-PLACE1000785	0.33
	PLACE1000786	F-PLACE1000786	0.88
40	PLACE1000793	F-PLACE1000793	0. 79
	PLACE1000798	F-PLACE1000798	0.59
	PLACE1000849	F-PLACE1000849	0.64
	PLACE1000856	F-PLACE1000856	0. 40
45	PLACE1000863	F-PLACE1000863	0.94
	PLACE1000909	F-PLACE1000909	0.94
	PLACE1000948	F-PLACE1000948	0.92
	PLACE1000972	F-PLACE1000972	0. 73
50	PLACE1000977	F-PLACE1000977	0.72
30	PLACE1000987	F-PLACE1000987	0. 93
	PLACE1001000	F-PLACE1001000	0.37
	PLACE1001010	F-PLACE1001010	0.50
	PLACE1001024	F-PLACE1001024	0.94
55	PLACE1001036	F-PLACE1001036	0.51

	PLACE1001054	F-PLACE1001054	0.88
	PLACE1001062	F-PLACE1001062	0.57
	PLACE1001092	F-PLACE1001092	0.94
5	PLACE1001104	F-PLACE1001104	0.38
	PLACE1001168	F-PLACE1001168	0.41
	PLACE1001171	F-PLACE1001171	0.36
	PLACE1001185	F-PLACE1001185	0, 93
10	PLACE1001238	F-PLACE1001238	0. 87
	PLACE1001241	F-PLACE1001241	0.67
	PLACE1001257	F-PLACE1001257	0.40
	PLACE1001280	F-PLACE1001280	0.70
15	PLACE1001294		0. 49
	PLACE1001304		
	PLACE1001311	F-PLACE1001311,	
	PLACE1001351	F-PLACE1001351	0.86
	PLACE1001366	F-PLACE1001366	0.39
20	PLACE1001383		0.94
	PLACE1001387	F-PLACE1001387	0.34
	PLACE1001399	F-PLACE1001399	0.38
	PLACE1001412	F-PLACE1001412	0.94
25	PLACE1001456	F-PLACE1001456	0.42
	PLACE1001484	F-PLACE1001484	0.88
	PLACE1001503	F-PLACE1001503	0.85
	PLACE1001545	F-PLACE1001545	0.92
30	PLACE1001551	F-PLACE1001551	0.39
	PLACE1001570	F-PLACE1001570	0.40
	PLACE1001602	F-PLACE1001602	0.60
	PLACE1001608	F-PLACE1001608	0.40
35	PLACE1001610	F-PLACE1001610	0.43
	PLACE1001632	F-PLACE1001632	0.48
	PLACE1001634	F-PLACE1001634	0.60
	PLACE1001692	F-PLACE1001692	0.74
40	PLACE1001729	F-PLACE1001729	0.69
	PLACE1001739	F-PLACE1001739	0.49
	PLACE1001740	F-PLACE1001740	0.50
	PLACE1001761	F-PLACE1001761	0.71
45	PLACE1001771	F-PLACE1001771	0.39
	PLACE1001781	F-PLACE1001781	0.57
	PLACE1001810	F-PLACE1001810	0. 57
	PLACE1001817	F-PLACE1001817	0.91
50	PLACE1001844	F-PLACE1001844	0. 34
50	PLACE1001869	F-PLACE1001869	0.34
	PLACE1001912	F-PLACE1001912	0.82
	PLACE1001920	F-PLACE1001920	0.85
	PLACE1001928	F-PLACE1001928	0.60
55	PLACE1001983	F-PLACE1001983	0.39

	PLACE1001989	F-PLACE1001989	0:74
	PLACE1002004	F-PLACE1002004	0.70
	PLACE1002046	F-PLACE1002046	0.81
5	PLACE1002072	F-PLACE1002072_	0. 94
	PLACE1002073	F-PLACE1002073	0.70
	PLACE1002090	F-PLACE1002090	0.64
	PLACE1002115	F-PLACE1002115	0.60
10	PLACE1002119	F-PLACE1002119	0.71
	PLACE1002140	F-PLACE1002140	0. 94
	PLACE1002163	F-PLACE1002163	0.56
	PLACE1002170	F-PLACE1002170	0.60
15	PLACE1002171	F-PLACE1002171	0.72
	PLACE1002213	F-PLACE1002213	0. 35
	PLACE1002342	F-PLACE1002342,	0.40
	PLACE1002395	F-PLACE1002395	0.87
20	PLACE1002433	F-PLACE1002433	0.73
20	PLACE1002437	F-PLACE1002437	0.46
	PLACE1002438	F-PLACE1002438	0.41
	PLACE1002450	F-PLACE1002450	0.48
	PLACE1002465	F-PLACE1002465	0.34
25	PLACE1002474	F-PLACE1002474	0.89
	PLACE1002499	F-PLACE1002499	0.94
	PLACE1002529	F-PLACE1002529	0.45
	PLACE1002532	F-PLACE1002532	0.76
30	PLACE1002571	F-PLACE1002571	0.40
	PLACE1002625	F-PLACE1002625	0.36
	PLACE1002655	F-PLACE1002655	0.41
	PLACE1002665	F-PLACE1002665	0.80
35	PLACE1002685	F-PLACE1002685	0. 31
	PLACE1002714	F-PLACE1002714	0. 90
	PLACE1002722	F-PLACE1002722	0. 43
	PLACE1002775		0.94
40	PLACE1002794		0. 90
	PLACE1002811		0.88
	PLACE1002815	F-PLACE1002815	0. 40
	PLACE1002816	F-PLACE1002816	0.73
45	PLACE1002834	F-PLACE1002834	0. 41
-	PLACE1002839	F-PLACE1002839	0. 73
	PLACE1002851	F-PLACE1002851	0.90
	PLACE1002908	F-PLACE1002908	0.94
50	PLACE1002941	F-PLACE1002941	0. 33
50	PLACE1002991	F-PLACE1002991	0.39
	PLACE1002993	F-PLACE1002993	0.94
	PLACE1002996	F-PLACE1002996	0.40
	PLACE1003025		0. 77
55	PLACE1003027	F-PLACE1003027	0. 54

	PLACE1003044	F-PLACE1003044	0.71
	PLACE1003045	F-PLACE1003045	0.89
	PLACE1003092	F-PLACE1003092	0.59
5	PLACE1003100	F-PLACE1003100	0.38
	PLACE1003108	F-PLACE1003108	0.94
	PLACE1003136	F-PLACE1003136	0. 87
	PLACE1003145	F-PLACE1003145	0.94
10	PLACE1003174	F-PLACE1003174	0.46
	PLACE1003176	F-PLACE1003176	0.48
	PLACE1003190	F-PLACE1003190	0. 57
	PLACE1003200	F-PLACE1003200	0.47
15	PLACE1003249	F-PLACE1003249	0. 33
13	PLACE1003256	F-PLACE1003256	0. 57
	PLACE1003296	F-PLACE1003296,	
	PLACE1003302	F-PLACE1003302	0. 94
	PLACE1003334	F-PLACE1003334	0. 78
20	PLACE1003342	F-PLACE1003342	0. 34
	PLACE1003353	F-PLACE1003353	0. 71
	PLACE1003366	F-PLACE1003366	0.66
	PLACE1003369	F-PLACE1003369	0. 91
25	PLACE1003383	F-PLACE1003383	0. 70
	PLACE1003394	F-PLACE1003394	0. 94
	PLACE1003454	F-PLACE1003454	0. 94
	PLACE1003493	F-PLACE1003493	0. 93
30	PLACE1003537	F-PLACE1003537	0. 94
	PLACE1003553	F-PLACE1003553	0.34
	PLACE1003592	F-PLACE1003592	0. 33
	PLACE1003596	F-PLACE1003596	0.89
35	PLACE1003602	F-PLACE1003602	0. 94
35	PLACE1003605	F-PLACE1003605	0. 94
	PLACE1003611	F-PLACE1003611	0.80
	PLACE1003625	F-PLACE1003625	0. 39
	PLACE1003669	F-PLACE1003669	0. 79
40	PLACE1003704	F-PLACE1003704	0. 43
	PLACE1003709	F-PLACE1003709	0. 90
	PLACE1003711	F-PLACE1003711	0.74
	PLACE1003723	F-PLACE1003723	0. 63
45	PLACE1003738	F-PLACE1003738	0. 91
	PLACE1003762	F-PLACE1003762	0.94
	PLACE1003771	F-PLACE1003771	0. 56
	PLACE1003783	F-PLACE1003783	0. 39
50	PLACE1003784	F-PLACE1003784	0.61
	PLACE1003795	F-PLACE1003795	0. 32
	PLACE1003870	F-PLACE1003870	0. 94
	PLACE1003885	F-PLACE1003885	0. 94
55	PLACE1003886	F-PLACE1003886	0. 94
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	PLACE1003888	F-PLACE1003888	0.36
	PLACE1003892	F-PLACE1003892	0. 92
	PLACE1003903	F-PLACE1003903	0. 68
5	PLACE1003915	F-PLACE1003915	
	PLACE1003923	F-PLACE1003923	0. 52
	PLACE1003936	F-PLACE1003936	0. 67
	PLACE1003968	F-PLACE1003968	0. 59
10	PLACE1003303	F-PLACE1004103	0. 94
	PLACE1004103	F-PLACE1004103	0. 44
	PLACE1004104	F-PLACE1004104	
			0. 91
45	PLACE1004128	F-PLACE1004128	0. 72
15	PLACE1004149		0. 41
	PLACE1004156	F-PLACE1004156	0. 50
	PLACE1004161	F-PLACE1004161,	
	PLACE1004183	F-PLACE1004183	0. 62
20	PLACE1004197	F-PLACE1004197	0. 39
	PLACE1004203	F-PLACE1004203	0. 94
	PLACE1004258	F-PLACE1004258	0. 46
	PLACE1004270	F-PLACE1004270	0. 65
25	PLACE1004277	F-PLACE1004277	0.86
	PLACE1004289	F-PLACE1004289	0. 79
	PLACE1004302	F-PLACE1004302	0. 57
	PLACE1004316	F-PLACE1004316	0.64
30	PLACE1004358	F-PLACE1004358	0. 85
30	PLACE1004376	F-PLACE1004376	0.64
	PLACE1004388	F-PLACE1004388	0.94
	PLACE1004405	F-PLACE1004405	0.82
	PLACE1004428	F-PLACE1004428	0. 68
35	PLACE1004437	F-PLACE1004437	0.94
	PLACE1004451	F-PLACE1004451	0. 39
	PLACE1004460	F-PLACE1004460	0. 92
	PLACE1004473	F-PLACE1004473	0.94
40	PLACE1004510	F-PLACE1004510	0. 94
	PLACE1004516	F-PLACE1004516	0.58
	PLACE1004548	F-PLACE1004548	0. 59
	PLACE1004550	F-PLACE1004550	0.94
45	PLACE1004564	F-PLACE1004564	0.53
10	PLACE1004629	F-PLACE1004629	0.34
	PLACE1004645	F-PLACE1004645	0.59
	PLACE1004646	F-PLACE1004646	0. 35
	PLACE1004664	F-PLACE1004664	0. 31
50	PLACE1004672	F-PLACE1004672	0.83
	PLACE1004674	F-PLACE1004674	0.90
	PLACE1004686	F-PLACE1004686	0. 31
	PLACE1004691	F-PLACE1004691	0.89
55	PLACE1004722	F-PLACE1004722	0. 48

	PLACE1004736	F-PLACE1004736	0.80
	PLACE1004740	F-PLACE1004740	0.34
	PLACE1004743	F-PLACE1004743	0. 70
5	PLACE1004751	F-PLACE1004751	0.49
	PLACE1004777	F-PLACE1004777	0. 72
	PLACE1004804	F-PLACE1004804	0. 46
	PLACE1004814	F-PLACE1004814	0. 70
10	PLACE1004824	F-PLACE1004824	0. 63
	PLACE1004868	F-PLACE1004868	0. 94
	PLACE1004885	F-PLACE1004885	0. 63
	PLACE1004902	F-PLACE1004902	0. 56
15	PLACE1004918	F-PLACE1004918	0. 85
	PLACE1004930	F-PLACE1004930	0. 83
	PLACE1004937	F-PLACE1004937	
	PLACE1004969	F-PLACE1004969	
	PLACE1004982	F-PLACE1004982	0. 61
20	PLACE1005026	F-PLACE1005026	0.81
	PLACE1005020	F-PLACE1005027	0. 91
	PLACE1005046	F-PLACE1005046	0. 31
	PLACE1005055	F-PLACE1005055	0. 57
25	PLACE1005055	F-PLACE1005066	0. 68
	PLACE1005000	F-PLACE1005101	0. 94
	PLACE1005101	F-PLACE1005101	0. 94
	PLACE1005102	F-PLACE1005102	0. 94
30	PLACE1005187	F-PLACE1005187	0. 94
	PLACE1005187	F-PLACE1005187	0. 67
	PLACE1005206	F-PLACE1005206	0. 72
	PLACE1005232	F-PLACE1005243	0. 72
35	PLACE1005243	F-PLACE1005261	0. 75
	PLACE1005261	F-PLACE1005266	0. 75
	PLACE1005277	F-PLACE1005277	0. 43
	PLACE1005277	F-PLACE1005287	0. 77
-	PLACE1005287	F-PLACE1005305	0. 77
40	PLACE1005308	F-PLACE1005308	0. 46
	PLACE1005313	F-PLACE1005313	0. 94
	PLACE1005313	F-PLACE1005313	0. 82
	PLACE1005327	F-PLACE1005327	0. 94
45	PLACE1005331	F-PLACE1005335	0. 94
	PLACE1005373	F-PLACE1005373	0. 71
	PLACE1005374	F-PLACE1005373	0. 71
	PLACE1005374 PLACE1005480	F-PLACE1005374	0. 82
50	PLACE1005480	F-PLACE1005480	0. 43
	PLACE1005481	F-PLACE1005481	0. 42
	PLACE1005494 PLACE1005530	F-PLACE1005530	0. 34
	PLACE1005550	F-PLACE1005530	0. 94
55	PLACE1005550	F-PLACE1005554	0. 86
	1 LAUL 1000004	F-FLAUE 1000004	v. 34

	PLACE1005557	F-PLACE1005557	0. 80
	PLACE1005595	F-PLACE1005595	0.69
_	PLACE1005603	F-PLACE1005603	0.92
5	PLACE1005623	F-PLACE1005623	0. 68
	PLACE1005630	F-PLACE1005630	0.58
	PLACE1005646	F-PLACE1005646	0.31
	PLACE1005656	F-PLACE1005656	0.85
10	PLACE1005698	F-PLACE1005698	0. 91
	PLACE1005727	F-PLACE1005727	0.94
	PLACE1005739	F-PLACE1005739	0. 87
	PLACE1005755	F-PLACE1005755	0. 32
15	PLACE1005763	F-PLACE1005763	0. 74
	PLACE1005799	F-PLACE1005799	0. 47
	PLACE1005803	F-PLACE1005803,	0.88
	PLACE1005804	F-PLACE1005804	0.36
20	PLACE1005813	F-PLACE1005813	0.76
	PLACE1005828	F-PLACE1005828	0.43
	PLACE1005834	F-PLACE1005834	0.89
	PLACE1005851	F-PLACE1005851	0.39
25	PLACE1005876	F-PLACE1005876	0.51
	PLACE1005890	F-PLACE1005890	0.81
	PLACE1005921	F-PLACE1005921	0. 90
	PLACE1005923	F-PLACE1005923	0.70
30	PLACE1005925	F-PLACE1005925	0.55
30	PLACE1005934	F-PLACE1005934	0. 94
	PLACE1005936	F-PLACE1005936	0. 41
	PLACE1005951	F-PLACE1005951	0.34
	PLACE1005953	F-PLACE1005953	0.31
35	PLACE1005955	F-PLACE1005955	0.89
	PLACE1005966	F-PLACE1005966	0. 73
	PLACE1005990	F-PLACE1005990	0.37
•	PLACE1006011	F-PLACE1006011	0.94
40	PLACE1006037	F-PLACE1006037	0.65
	PLACE1006040	F-PLACE1006040	0.94
	PLACE1006119		
	PLACE1006139	F-PLACE1006139	0.94
45	PLACE1006157	F-PLACE1006157	0.34
	PLACE1006159	F-PLACE1006159	0.50
	PLACE1006167	F-PLACE1006167	0.94
	PLACE1006170	F-PLACE1006170	0.94
50	PLACE1006195	F-PLACE1006195	0.31
	PLACE1006196	F-PLACE1006196 F-PLACE1006225	0.84
	PLACE1006225		0.87
	PLACE1006236	F-PLACE1006236 F-PLACE1006239	0. 92 0. 53
55	PLACE1006239	F-PLACE1006239	0. 53
	PLACE1006246	F-FLAUCIUU0240	u. <del>44</del>

	PLACE1006248	F-PLACE1006248	0.49
	PLACE1006325	F-PLACE1006325	0.74
	PLACE1006335	F-PLACE1006335	0.82
5	PLACE1006357	F-PLACE1006357	0. 33
	PLACE1006385	F-PLACE1006385	0. 54
	PLACE1006412	F-PLACE1006412	0. 79
	PLACE1006414	F-PLACE1006414	0. 52
10	PLACE1006438	F-PLACE1006438	0. 94
	PLACE1006445	F-PLACE1006445	0. 37
	PLACE1006469	F-PLACE1006469	0. 85
	PLACE1006482	F-PLACE1006482	0. 94
15	PLACE1006488	F-PLACE1006488	0. 94
	PLACE1006492	F-PLACE1006492	0. 32
	PLACE1006531	F-PLACE1006531,	0. 45
	PLACE1006552	F-PLACE1006552	0. 42
20	PLACE1006598	F-PLACE1006598	0. 33
	PLACE1006615	F-PLACE1006615	0.89
	PLACE1006626	F-PLACE1006626	0.69
	PLACE1006673	F-PLACE1006673	0.61
25	PLACE1006678	F-PLACE1006678	0.53
25	PLACE1006704	F-PLACE1006704	0. 47
	PLACE1006731	F-PLACE1006731	0.72
	PLACE1006782	F-PLACE1006782	0.68
•	PLACE1006819	F-PLACE1006819	0. 57
30	PLACE1006829	F-PLACE1006829	0. 47
	PLACE1006883	F-PLACE1006883	0.88
	PLACE1006901	F-PLACE1006901	0. 43
	PLACE1006917	F-PLACE1006917	0.56
35	PLACE1006932	F-PLACE1006932	0. 45
	PLACE1006935	F-PLACE1006935	0.58
	PLACE1006956	F-PLACE1006956	0.77
	PLACE1006958	F-PLACE1006958	0. 75
40	PLACE1006961	F-PLACE1006961	0.34
	PLACE1006962	F-PLACE1006962	0.94
	PLACE1006966	F-PLACE1006966	0.56
	PLACE1006989	F-PLACE1006989	0. 47
45	PLACE1007014	F-PLACE1007014	0.94
	PLACE1007021 PLACE1007053	F-PLACE1007021	0.74
	PLACE1007053	F-PLACE1007053	0.73
	PLACE100708	F-PLACE1007068 F-PLACE1007105	0. 43
50	PLACE1007105	F-PLACE1007105	0. 88 0. 82
	PLACE1007178	F-PLACE1007178	0. 82
	PLACE1007178	F-PLACE1007178	0. 79
	PLACE1007228	F-PLACE1007238	0.79
55	PLACE1007239	F-PLACE1007239	0. 94
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	· PLACE1007242	F-PLACE1007242	0.48
	PLACE1007243	F-PLACE1007243	0.77
•	PLACE1007257	F-PLACE1007257	0. 54
5	PLACE1,00727,4	F-PLACE1007274_	0.94
	PLACE1007282	F-PLACE1007282	0.94
	PLACE1007301	F-PLACE1007301	0. 34
	PLACE1007317	F-PLACE1007317	0. 43
10	PLACE1007342	F-PLACE1007342	0. 72
	PLACE1007346	F-PLACE1007346	0.92
	PLACE1007367	F-PLACE1007367	0. 43
	PLACE1007375	F-PLACE1007375	0. 85
15	PLACE1007386	F-PLACE1007386	0.58
	PLACE1007402	F-PLACE1007402	0. 74
	PLACE1007409	F-PLACE1007409,	
	PLACE1007416	F-PLACE1007416	0. 80
20	PLACE1007450	F-PLACE1007450	0.41
	PLACE1007452	F-PLACE1007452	0. 68
	PLACE1007454	F-PLACE1007454	0.39
	PLACE1007460	F-PLACE1007460	0.58
25	PLACE1007484	F-PLACE1007484	0.94
	PLACE1007488	F-PLACE1007488	0.50
	PLACE1007507		0.44
	PLACE1007511	F-PLACE1007511	0. 86 0. 66
30	PLACE1007524 PLACE1007537	F-PLACE1007524 F-PLACE1007537	0. 93
	PLACE1007537 PLACE1007544	F-PLACE1007537	0. 93
	PLACE1007544 PLACE1007547	F-PLACE1007547	0. 65
	PLACE1007547 PLACE1007583	F-PLACE1007583	0. 87
35	PLACE1007598	F-PLACE1007598	0.39
	PLACE1007618	F-PLACE1007618	0.62
	PLACE1007621	F-PLACE1007621	0. 63
	PLACE1007632	F-PLACE1007632	0. 75
40	PLACE1007645	F-PLACE1007645	0. 94
	PLACE1007649	F-PLACE1007649	0. 56
	PLACE1007688	F-PLACE1007688	0.94
	PLACE1007690	F-PLACE1007690	0. 44
45	PLACE1007697	F-PLACE1007697	0. 93
45	PLACE1007705	F-PLACE1007705	0.61
	PLACE1007706	F-PLACE1007706	0.47
	PLACE1007725	F-PLACE1007725	0.61
	PLACE1007729	F-PLACE1007729	0. 53
50	PLACE1007730	F-PLACE1007730	0.31
	PLACE1007746	F-PLACE1007746	0.81
	PLACE1007791	F-PLACE1007791	0.46
	PLACE1007810	F-PLACE1007810	0. 73
55	PLACE1007843	F-PLACE1007843	0. 31

		PLACE1007846	F-PLACE1007846	0. 38
		PLACE1007858	F-PLACE1007858	0.79
		PLACE1007897	F-PLACE1007897	0.94
5		PLACE1007908	F-PLACE1007908	0.94
		PLACE1007946	F-PLACE1007946	0. 52
		PLACE1007954	F-PLACE1007954	0.43
		PLACE1007955	F-PLACE1007955	0. 41
10	)	PLACE1007958	F-PLACE1007958	0.37
		PLACE1007969	F-PLACE1007969	0.87
		PLACE1007990	F-PLACE1007990	0.94
		PLACE1008000	F-PLACE1008000	0.94
1:	5	PLACE1008002	F-PLACE1008002	0.93
		PLACE1008044	F-PLACE1008044	0.72
		PLACE1008095	F-PLACE1008095,	0.94
		PLACE1008122	F-PLACE1008122	0.94
20	1	PLACE1008129	F-PLACE1008129	0.80
		PLACE1008132	F-PLACE1008132	0. 37
			F-PLACE1008177	0.74
		PLACE1008198		0. 94
25	i	PLACE1008209		0.54
		PLACE1008273	F-PLACE1008273	0.94
		PLACE1008275		0.82
		PLACE1008280		0.34
30	1	PLACE1008309		0. 90
		PLACE1008329		0.38
		PLACE1008356	F-PLACE1008356	0.93
		PLACE1008368	F-PLACE1008368	0.43
35		PLACE1008398	F-PLACE1008398	0.78
3.		PLACE1008401 PLACE1008402	F-PLACE1008401 F-PLACE1008402	0. 81 0. 88
		PLACE1008402		0. 39
		PLACE1008429		0. 39
.,		PLACE1008465		0. 77
40		PLACE1008488		0. 42
		PLACE1008524	F-PLACE1008524	0.41
		PLACE1008531	F-PLACE1008531	0.94
		PLACE1008532	F-PLACE1008532	0.61
45		PLACE1008533	F-PLACE1008533	0. 45
		PLACE1008568	F-PLACE1008568	0.65
		PLACE1008603	F-PLACE1008603	0.42
		PLACE1008626	F-PLACE1008626	0. 94
50		PLACE1008627	F-PLACE1008627	0.72
		PLACE1008629	F-PLACE1008629	0.33
		PLACE1008650	F-PLACE1008650	0.79
		PLACE1008693	F-PLACE1008693	0. 32
55		PLACE1008696	F-PLACE1008696	0.39

		PLACE1008790	F-PLACE1008790	0. 81
		PLACE1008808	F-PLACE1008808	0.51
		PLACE1008813		0.36
	5	PLACE1008854		
		PLACE1008867	F-PLACE1008867	0. 34
		PLACE1008887	F-PLACE1008887	0.49
		PLACE1008902	F-PLACE1008902	0.91
	10	PLACE1008925	F-PLACE1008925	0.94
		PLACE1008934		0.70
		PLACE1009027		0. 32
		PLACE1009045	F-PLACE1009045	0.42
	15	PLACE1009060	F-PLACE1009060	0. 69
		PLACE1009090	F-PLACE1009090	0. 46
		PLACE1009091	F-PLACE1009091,	
		PLACE1009094	•	
	20		F-PLACE1009099	
	20		F-PLACE1009110	
			F-PLACE1009111	0.80
			F-PLACE1009113	0. 88
		PLACE1009130	F-PLACE1009130	0.60
	25	PLACE1009158	F-PLACE1009158	0.41
		PLACE1009166	F-PLACE1009166	0.94
		PLACE1009174	F-PLACE1009174	0.38
		PLACE1009186	F-PLACE1009186	0.59
:	30	PLACE1009190	F-PLACE1009190	0.66
		PLACE1009230	F-PLACE1009230	0.87
		PLACE1009298	F-PLACE1009298	0.41
		PLACE1009319	F-PLACE1009319	0.53
;	35	PLACE1009328	F-PLACE1009328	0.66
		PLACE1009335	F-PLACE1009335	0.47
		PLACE1009338	F-PLACE1009338	0.70
		PLACE1009368	F-PLACE1009368	0.94
4	10	PLACE1009375	F-PLACE1009375	0.76
		PLACE1009388	F-PLACE1009388	0.46
		PLACE1009404	F-PLACE1009404	0.72
	,	PLACE1009434	F-PLACE1009434	0.50
4	15	PLACE1009443	F-PLACE1009443	0.90
		PLACE1009444	F-PLACE1009444	0.33
		PLACE1009459	F-PLACE1009459	0.94
		PLACE1009468	F-PLACE1009468	0.88
	50	PLACE1009476	F-PLACE1009476	0.84
	,	PLACE1009524	F-PLACE1009524	0.88
		PLACE1009542	F-PLACE1009542	0.55
		PLACE1009571	F-PLACE1009571	0.61
		PLACE1009581	F-PLACE1009581	0.94
5	55	PLACE1009596	F-PLACE1009596	0.44

	PLACE1009607	F-PLACE1009607	0. 47
	PLACE1009622	F-PLACE1009622	0.65
	PLACE1009659	F-PLACE1009659	0.36
5	PLACE1009665	F-PLACE1009665	0.43
	PLACE1009670	F-PLACE1009670	0. 33
	PLACE1009708	F-PLACE1009708	0. 80
	PLACE1009721	F-PLACE1009721	0.81
10	PLACE1009731	F-PLACE1009731	0.47
	PLACE1009763	F-PLACE1009763	0. 94
	PLACE1009794	F-PLACE1009794	0.33
	PLACE1009798	F-PLACE1009798	0.34
15	PLACE1009845	F-PLACE1009845	0.66
	PLACE1009861	F-PLACE1009861	0. 72
	PLACE1009886	F-PLACE1009886,	0.63
	PLACE1009908	F-PLACE1009908	0.92
20	PLACE1009921	F-PLACE1009921	0.70
-	PLACE1009971	F-PLACE1009971	0.40
	PLACE1009992	F-PLACE1009992	0. 89
	PLACE1009995	F-PLACE1009995	0.88
25	PLACE1009997	F-PLACE1009997	0.41
-	PLACE1010023		0.90
	PLACE1010031	F-PLACE1010031	0. 71
		F-PLACE1010053	0.51
30	PLACE1010074		0. 75
30	PLACE1010076		0. 53
	PLACE1010096		0. 44
	PLACE1010102	F-PLACE1010102	0.77
05	PLACE1010105 PLACE1010106	F-PLACE1010105 F-PLACE1010106	0. 58 0. 39
35	PLACE1010108		0. 85
	PLACE1010134		0. 32
	PLACE1010152		0. 94
_	PLACE1010194		0.94
40	PLACE1010202	F-PLACE1010202	0. 33
	PLACE1010261		
	PLACE1010274	F-PLACE1010274	0.94
	PLACE1010293	F-PLACE1010293	0.55
45	PLACE1010310	F-PLACE1010310	0. 41
	PLACE1010321	F-PLACE1010321	0.40
	PLACE1010324	F-PLACE1010324	0.40
	PLACE1010329	F-PLACE1010329	0.36
50	PLACE1010362	F-PLACE1010362	0.34
	PLACE1010364	F-PLACE1010364	0.37
	PLACE1010383	F-PLACE1010383	0.94
	PLACE1010401	F-PLACE1010401	0.53
55	PLACE1010481	F-PLACE1010481	0.74

	PLACE1010491	F-PLACE1010491	0.31
	PLACE1010492	F-PLACE1010492	0.39
	PLACE1010522	F-PLACE1010522	0.92
5	PLACE1010529	F-PLACE1010529	0. 65
	PLACE1010547	F-PLACE1010547	0.82
	PLACE1010580	F-PLACE1010580	0.72
	PLACE1010599	F-PLACE1010599	0. 83
10	PLACE1010616	F-PLACE1010616	0. 54
	PLACE1010622	F-PLACE1010622	0, 92
	PLACE1010629	F-PLACE1010629	0. 94
	PLACE1010630	F-PLACE1010630	0. 37
15	PLACE1010661	F-PLACE1010661	0.56
13	PLACE1010714	F-PLACE1010714	0.68
	PLACE1010714	F-PLACE1010720,	
	PLACE1010723	F-PLACE1010743	0. 75
	PLACE1010743	F-PLACE1010761	0. 73
20	PLACE1010701	F-PLACE1010771	0. 94
	PLACE1010771	F-PLACE1010771	0.80
	PLACE1010800	F-PLACE1010800	0. 41
	PLACE1010800	F-PLACE1010800	0. 94
25	PLACE1010811	F-PLACE1010833	0. 94
	PLACE1010835 PLACE1010856	F-PLACE1010856	0. 37
	PLACE1010857	F-PLACE1010857	0. 54
	PLACE1010857	F-PLACE1010870	0. 50
30	PLACE1010870 PLACE1010877	F-PLACE1010877	0. 50
	PLACE1010900	F-PLACE1010900	0.77
	PLACE1010900 PLACE1010917	F-PLACE1010900	
	PLACE1010917 PLACE1010925	F-PLACE1010917	0. 94 0. 79
0.5	PLACE1010925 PLACE1010926	F-PLACE1010925	0.79
35	PLACE1010920 PLACE1010942	F-PLACE1010920	0. 36
	PLACE1010942	F-PLACE1010942	0. 61
	PLACE1010944 PLACE1010954	F-PLACE1010944	0.83
	PLACE1010954 PLACE1010960	F-PLACE1010954	0. 83
40	PLACE1010960 PLACE1010965	F-PLACE1010965	0. 68
	PLACE1011026	F-PLACE1010905	0. 38
	PLACE1011026 PLACE1011041		
	PLACE1011041 PLACE1011046	F-PLACE1011041 F-PLACE1011046	0. 94 0. 85
45	PLACE1011046 PLACE1011054		
	PLACE1011054 PLACE1011056	F-PLACE1011054 F-PLACE1011056	0.65
			0.82
	PLACE1011090	F-PLACE1011090	0.94
50	PLACE1011114	F-PLACE1011114	0. 94
	PLACE1011160	F-PLACE1011160	0. 62
	PLACE1011214	F-PLACE1011214	0.40
	PLACE1011219	F-PLACE1011219	0.48
	PLACE1011221	F-PLACE1011221	0.94
55	PLACE1011229	F-PLACE1011229	0. 46

	DI ACE1011000	E DI 1051011000	0 04
	PLACE1011263	F-PLACE1011263	0.94
	PLACE1011273	F-PLACE1011273	0. 32
5	PLACE1011291	F-PLACE1011291	0.55
,	PLACE1011310	F-PLACE1011310_	• • • •
	PLACE1011325	F-PLACE1011325	0.39
	PLACE1011332	F-PLACE1011332	0.74
	PLACE1011371	F-PLACE1011371	0.32
10	PLACE1011399	F-PLACE1011399	0.89
	PLACE1011433	F-PLACE1011433	0. 33
	PLACE1011472	F-PLACE1011472	0.59
	PLACE1011477	F-PLACE1011477	0.80
15	PLACE1011563	F-PLACE1011563	0.83
	PLACE1011586	F-PLACE1011586	0.82
	PLACE1011635	F-PLACE1011635,	0.62
	PLACE1011646	F-PLACE1011646	0.37
20	PLACE1011664	F-PLACE1011664	0.65
	PLACE1011675	F-PLACE1011675	0.89
	PLACE1011682	F-PLACE1011682	0.81
	PLACE1011725	F-PLACE1011725	0.86
25	PLACE1011783	F-PLACE1011783	0.78
25	PLACE1011858	F-PLACE1011858	0.76
	PLACE1011875	F-PLACE1011875	0.49
	PLACE1011891	F-PLACE1011891	0.34
	PLACE1011896	F-PLACE1011896	0.34
30	PLACE1011922	F-PLACE1011922	0.46
	PLACE1011923	F-PLACE1011923	0.74
	PLACE1011982	F-PLACE1011982	0.71
	PLACE1012031	F-PLACE1012031	0.80
35	PLACE2000006	F-PLACE2000006	0.61
	PLACE2000007	F-PLACE2000007	0.37
	PLACE2000014	F-PLACE2000014	0.67
	PLACE2000015	F-PLACE2000015	0.53
40	PLACE2000021	F-PLACE2000021	0.36
	PLACE2000030	F-PLACE2000030	0.83
	PLACE2000033	F-PLACE2000033	0.44
	PLACE2000034	F-PLACE2000034	0.88
45	PLACE2000039	F-PLACE2000039	0.67
	PLACE2000047	F-PLACE2000047	0.84
	PLACE2000050	F-PLACE2000050	0.41
	PLACE2000062	F-PLACE2000062	0.94
	PLACE2000072	F-PLACE2000072	0.86
50	PLACE2000097	F-PLACE2000097	0.94
	PLACE2000100	F-PLACE2000100	0.75
	PLACE2000111	F-PLACE2000111	0.62
	PLACE2000124	F-PLACE2000124	0.77
55	PLACE2000164	F-PLACE2000164	0.57

	•		
	PLACE2000170	F~PLACE2000170	0.60
	PLACE2000172	F-PLACE2000172	0. 55
	PLACE2000223	F-PLACE2000223	0.54
5	PLACE2000235	F-PLACE2000235	0. 94
	PLACE2000246	F-PLACE2000246	0.94
	PLACE2000274	F-PLACE2000274	0.82
	PLACE2000302	F-PLACE2000302	0. 92
10	PLACE2000341	F-PLACE2000341	0. 58
	PLACE2000359	F-PLACE2000359	0, 33
	PLACE2000371	F-PLACE2000371	0.71
	PLACE2000373	F-PLACE2000373	0. 65
15	PLACE2000399	F-PLACE2000399	0.41
,,	PLACE2000404		0. 94
	PLACE2000404	F-PLACE2000404	
	PLACE2000477	F-PLACE2000427	0.60
	PLACE2000427	F-PLACE2000427	0.61
20	PLACE2000438	F-PLACE2000438	0. 56
	PLACE2000458	F-PLACE2000458	0. 82
	PLACE3000009	F-PLACE3000009	0. 56
	PLACE3000020	F-PLACE3000020	0. 94
25	PLACE3000029	F-PLACE3000029	0. 94
	PLACE3000059	F-PLACE3000059	0. 51
	PLACE3000070	F-PLACE3000070	0. 67
	PLACE3000103	F-PLACE3000103	0. 34
30	PLACE3000121	F-PLACE3000121	0. 32
	PLACE3000142	F-PLACE3000142	0. 67
	PLACE3000145	F-PLACE3000145	0.84
	PLACE3000147	F-PLACE3000147	0.57
35	PLACE3000149	F-PLACE3000148	0. 94
	PLACE3000155	F-PLACE3000155	0. 42
	PLACE3000156	F-PLACE3000156	0. 40
	PLACE3000157	F-PLACE3000157	0. 75
40	PLACE3000160	F-PLACE3000160	0. 44
40	PLACE3000194	F-PLACE3000194	0. 37
	PLACE3000197	F-PLACE3000197	0. 37
	PLACE3000221	F-PLACE3000221	0. 49
	PLACE3000226	F-PLACE3000226	0. 39
45	PLACE3000242	F-PLACE3000242	0. 94
	PLACE3000244	F-PLACE3000244	0. 56
	PLACE3000254	F-PLACE3000254	0.60
	PLACE3000304	F-PLACE3000304	0. 42
50	PLACE3000310	F-PLACE3000310	0. 71
	PLACE3000320	F-PLACE3000320	0. 37
	PLACE3000322	F-PLACE3000322	0. 46
	PLACE3000331	F-PLACE3000331	0. 31
55	PLACE3000339	F-PLACE3000339	0. 45

	PLACE3000350	F-PLACE3000350	0. 52 <sup>-</sup>
	PLACE3000352	F-PLACE3000352	0. 65
	PLACE3000353	F-PLACE3000353	0.71°
5	PLACE3000363	F-PLACE3000363	0.45
	PLACE3000405	F-PLACE3000405	0.52
	PLACE3000413	F-PLACE3000413	0.85
	PLACE3000416	F-PLACE3000416	0.94
10	PLACE3000425	F-PLACE3000425	0.40
	PLACE3000455	F-PLACE3000455	0.45
	PLACE3000477	F-PLACE3000477	0.79
	PLACE4000009	F-PLACE4000009	0.74
15	PLACE4000014	F-PLACE4000014	0.46
	PLACE4000034	F-PLACE4000034	0.89
	PLACE4000049	F-PLACE4000049,	0. 68
	PLACE4000052	F-PLACE4000052	0.80
20	PLACE4000063	F-PLACE4000063	0.69
	PLACE4000089	F-PLACE4000089	0.44
	PLACE4000106	F-PLACE4000106	0. 34
	PLACE4000128	F-PLACE4000128	0. 89
25	PLACE4000129	F-PLACE4000129	0. 61
	PLACE4000131	F-PLACE4000131	0. 55
	PLACE4000147	F-PLACE4000147	0. 94
	PLACE4000156	F-PLACE4000156	0. 37
30	PLACE4000192	F-PLACE4000192	0.88
30	PLACE4000211	F-PLACE4000211	0.64
	PLACE4000222	F-PLACE4000222	0.40
	PLACE4000230	F-PLACE4000230	0.49
	PLACE4000259	F-PLACE4000259	0.84
35	PLACE4000261	F-PLACE4000261	0. 42
	PLACE4000269	F-PLACE4000269	0.60
	PLACE4000270	F-PLACE4000270	0.61
	PLACE4000300	F-PLACE4000300	0. 85
40	PLACE4000323	F-PLACE4000323	0. 76
	PLACE4000326	F-PLACE4000326	0.86
	PLACE4000369	F-PLACE4000369	0. 87
	PLACE4000387	F-PLACE4000387	0. 41
45	PLACE4000392	F-PLACE4000392	0.39
	PLACE4000431	F-PLACE4000431	0.88
	PLACE4000445	F-PLACE4000445	0.40
	PLACE4000450	F-PLACE4000450	0.89
50	PLACE4000465	F-PLACE4000465	0. 38 0. 44
	PLACE4000487 PLACE4000489	F-PLACE4000487 F-PLACE4000489	0. 44
	PLACE4000489 PLACE4000521	F-PLACE4000489	0.38
	PLACE4000521 PLACE4000522	F-PLACE4000521	0. 38
55	PLACE4000522 PLACE4000558	F-PLACE4000522	
	PLAUE4000558	r-rlage4000558	0. 44

	PLACE4000581	F-PLACE4000581	0. 59
	PLACE4000590	F-PLACE4000590	0.38
	PLACE4000593	F-PLACE4000593	0.33
5	PLACE4000612	F-PLACE4000612	0.37
	PLACE4000638	F-PLACE4000638	0. 38
	PLACE4000650	F-PLACE4000650	0. 59
	PLACE4000654	F-PLACE4000654	0. 57
10	PLACE4000670	F-PLACE4000670	0.71
	SKNMC1000011	F-SKNMC1000011	0.94
	SKNMC1000013	F-SKNMC1000013	0.66
	SKNMC1000046	F-SKNMC1000046	0.72
15	SKNMC1000050	F-SKNMC1000050	0.47
	SKNMC1000091	F-SKNMC1000091	0.68
	THYR01000017	F-THYR01000017,	0. 93
	THYR01000034	F-THYR01000034	0. 48
20	THYR01000040	F-THYR01000040	0. 32
	THYR01000070	F-THYR01000070	0. 75
	THYR01000072	F-THYR01000072	0. 94
	THYR01000107	F-THYR01000107	0. 94
25	THYR01000121	F-THYR01000121	0.46
25	THYR01000124	F-THYR01000124	0.37
	THYR01000163	F-THYR01000163	0.45
	THYR01000173	F-THYR01000173	0.94
	THYR01000197	F-THYR01000197	0.78
30	THYR01000199	F-THYR01000199	0. 55
	THYRO1000206	F-THYR01000206	0.59
	THYR01000242	F-THYR01000242	0.69
	THYRO1000253	F-THYR01000253	0.48
35	THYR01000270	F-THYR01000270	0.85
	THYRO1000288	F-THYR01000288	0.31
	THYR01000320	F-THYR01000320	0.94
	THYR01000327	F-THYR01000327	0.44
40	THYRO1000343	F-THYR01000343	0.82
	THYR01000358	F-THYR01000358	0. 89
	THYR01000368	F-THYR01000368	0. 94
	THYR01000381	F-THYR01000381	0. 64
45	THYR01000387	F-THYR01000387	0. 57
	THYR01000394	F-THYR01000394	0. 94
	THYR01000395	F-THYR01000395	0. 91
	THYR01000401	F-THYR01000401	0. 73
50	THYR01000488	F-THYR01000488	0. 94
<b>50</b>	THYR01000501	F-THYR01000501	0. 74
	THYR01000502	F-THYR01000502	0. 64
	THYR01000505	F-THYR01000505	0. 47
	THYR01000569	F-THYR01000569	0. 76
55	THYR01000570	F-THYR01000570	0. 79

	THYR01000585	F-THYR01000585	0.60
	THYR01000605	F-THYR01000605	0. 79
	THYR01000637	F-THYR01000637	0.80
5	THYR01000662	F-THYR01000662	
	THYR01000666	F-THYR01000666	0.47
	THYR01000676	F-THYR01000676	0. 94
	THYR01000684	F-THYR01000684	0. 72
10	**	F-THYR01000084	0.72
	THYR01000712 THYR01000715	F-THYR01000715	0. 94
		F-THYR01000713	0. 44
	THYR01000734	F-THYR01000756	0. 94
45	THYR01000756	F-THYR01000756	0. 73
15	THYR01000777		0. 73
	THYR01000783	F-THYR01000783	0. 74
	THYR01000787	F-THYR01000787, F-THYR01000829	0. 43
	THYR01000829	F-THYR01000829	0. 43
20	THYR01000855 THYR01000916	F-THYR01000855	0. 64
	THYR01000916	F-THYR01000916	0. 39
	THYR01000926	F-THYR01000926	0. 94
	THYR01000951	F-THYR01000951	0. 94
25	THYR01000952	F-THYR01000952	0. 67
	THYR01000983	F-THYR01000983	0. 87
	THYR01000984	F-THYR01000984	0. 94
	THYR01000988	F-THYR01000988	0. 37
30	THYR01001033	F-THYR01001033	0.74
	THYR01001100	F-THYR01001100	0.74
	THYR01001120	F-THYR01001120	0. 87
	THYR01001134	F-THYR01001134	0. 93
35	THYR01001142	F-THYR01001142	0. 61
	THYR01001173	F-THYR01001173	0.50
	THYR01001189	F-THYR01001189	0.40
	THYR01001204	F-THYR01001204	0. 94
40	THYR01001213	F-THYR01001213	0.94
	THYR01001271	F-THYR01001271	0.93
	THYR01001287	F-THYR01001287	0.32
	THYR01001313	F-THYR01001313	0.53
45	THYR01001320	F-THYR01001320	0.68
40	THYR01001347	F-THYR01001347	0.68
	THYR01001363	F-THYR01001363	0.39
	THYR01001374	F-THYR01001374	0.94
	THYR01001403	F-THYR01001403	0.49
50	THYR01001405	F-THYR01001405	0. 90
	THYR01001406	F-THYR01001406	0.67
	THYR01001458	F-THYR01001458	0. 94
	THYR01001487	F-THYR01001487	0. 63
55	THYR01001537	F-THYR01001537	0. 94

	THYR01001584	F-THYR01001584	0.77
	THYR01001602	F-THYR01001602	0.64
	THYR01001605	F-THYR01001605	0.52
5	THYR01001637	F-THYR01001637_	0. 68
	THYR01001656	F-THYR01001656	0.82
	THYR01001661	F-THYR01001661	0.94
	THYR01001671	F-THYR01001671	0.94
10	THYR01001703	F-THYR01001703	0.87
	THYR01001721	F-THYR01001721	0.38
	THYR01001738	F-THYR01001738	0.51
	THYR01001793	F-THYR01001793	0.84
15	THYR01001809	F-THYR01001809	0.73
15	THYR01001828	F-THYR01001828	0.88
	THYR01001854	F-THYR01001854	
	VESEN1000122	F-VESEN1000122	0.36
	Y79AA1000013	F-Y79AA1000013	0.94
20	Y79AA1000037	F-Y79AA1000037	0.94
	Y79AA1000059	F-Y79AA1000059	0.94
	Y79AA1000131	F-Y79AA1000131	0. 56
	Y79AA1000181	F-Y79AA1000181	0. 59
25	Y79AA1000202	F-Y79AA1000202	0, 45
	Y79AA1000214	F-Y79AA1000214	0. 93
	Y79AA1000230	F-Y79AA1000230	0.94
	Y79AA1000231	F-Y79AA1000231	0.36
30	Y79AA1000258	F-Y79AA1000258	0.36
	Y79AA1000268	F-Y79AA1000268	0.92
	Y79AA1000313	F-Y79AA1000313	0.91
	Y79AA1000328	F-Y79AA1000328	0.94
35	Y79AA1000342	F-Y79AA1000342	0.83
	Y79AA1000349	F-Y79AA1000349	0.50
	Y79AA1000355	F-Y79AA1000355	0.94
	Y79AA1000368	F-Y79AA1000368	0. 91
40	Y79AA1000420	F-Y79AA1000420	0.94
40	Y79AA1000469	F-Y79AA1000469	0.51
	Y79AA1000480	F-Y79AA1000480	0.94
	Y79AA1000539	F-Y79AA1000539	0.74
	Y79AA1000540	F-Y79AA1000540	0.94
45	Y79AA1000560	F-Y79AA1000560	0.93
	Y79AA1000574	F-Y79AA1000574	0.87
	Y79AA1000589	F-Y79AA1000589	0.94
	Y79AA1000627	F-Y79AA1000627	0.55
50	Y79AA1000705	F-Y79AA1000705	0.76
	Y79AA1000734	F-Y79AA1000734	0.60
	Y79AA1000748	F-Y79AA1000748	0.45
	Y79AA1000752	F-Y79AA1000752	0.94
55	Y79AA1000774	F-Y79AA1000774	0.38

	Y79AA1000782 F-Y79AA1000782	0.47
	Y79AA1000784 F-Y79AA1000784	0. 93
-	Y79AA1000794 F-Y79AA1000794	0.67
5	Y79AA1000800 F-Y79AA1000800	
	Y79AA1000824 F-Y79AA1000824	0.90
	Y79AA1000827 F-Y79AA1000827	0.94
	Y79AA1000833 F-Y79AA1000833	0.34
10	Y79AA1000850 F-Y79AA1000850	0. 92
	Y79AA1000962 F-Y79AA1000962	0. 69
	Y79AA1000966 F-Y79AA1000966	0. 64
	Y79AA1000968 F-Y79AA1000968	0.64
15	Y79AA1000976 F-Y79AA1000976	0. 66
	Y79AA1000985 F-Y79AA1000985	0.66
	Y79AA1001023 F-Y79AA1001023 a	0. 33
	Y79AA1001041 F-Y79AA1001041	0.90
20	Y79AA1001048 F-Y79AA1001048	0.81
	Y79AA1001068 F-Y79AA1001068	0.64
	Y79AA1001077 F-Y79AA1001077	0.78
	Y79AA1001078 F-Y79AA1001078	0.92
25	Y79AA1001105 F-Y79AA1001105	0.70
	Y79AA1001145 F-Y79AA1001145	0.46
	Y79AA1001177 F-Y79AA1001177	0.74
	Y79AA1001185 F~Y79AA1001185	0. 74
30	Y79AA1001211 F-Y79AA1001211	0. 94
30	Y79AA1001216 F-Y79AA1001216	0.61
	Y79AA1001228 F-Y79AA1001228	0. 56
	Y79AA1001233 F~Y79AA1001233	0. 83
	Y79AA1001236 F-Y79AA1001236	0.32
35	Y79AA1001281 F-Y79AA1001281	0.51
	Y79AA1001299 F-Y79AA1001299	0.94
	Y79AA1001312 F-Y79AA1001312	0. 53
	Y79AA1001323 F-Y79AA1001323	0.76
40	Y79AA1001391 F-Y79AA1001391	0.83
	Y79AA1001394 F-Y79AA1001394	0.94
	Y79AA1001402 F-Y79AA1001402	0.90
	Y79AA1001493 F-Y79AA1001493 Y79AA1001511 F-Y79AA1001511	0.94
45		0.47
		0.92
	•	0.32
		0.50
50	Y79AA1001555 F-Y79AA1001555 Y79AA1001581 F-Y79AA1001581	0.92
	Y79AA1001585 F-Y79AA1001585	0.94
	Y79AA1001603 F-Y79AA1001603	0. 80 0. 94
	Y79AA1001603 F-Y79AA1001613	0. 94
55	Y79AA1001613 F-179AA1001613	0. 49
	112001001041 1 113001001041	J. 73

	Y79AA1001665	F-Y79AA1001665	0. 78
	Y79AA1001679	F-Y79AA1001679	0. 94
	Y79AA1001696	F-Y79AA1001696	0. 60
5	Y79AA1001705	F-Y79AA1001705_	0. 67
	Y79AA1001711	F-Y79AA1001711	0. 59
	Y79AA1001781	F-Y79AA1001781	0. 74
10	Y79AA1001805	F-Y79AA1001805	0. 79
10	Y79AA1001827	F-Y79AA1001827	0.84
	Y79AA1001846	F-Y79AA1001846	0.34
	Y79AA1001866	F-Y79AA1001866	0. 75
15	Y79AA1001874	F-Y79AA1001874	0.59
	Y79AA1001923	F-Y79AA1001923	0.32
	Y79AA1001963	F-Y79AA1001963	0.51
	Y79AA1002027	F-Y79AA1002027	0.39
20	Y79AA1002083	F-Y79AA1002083	0.65
	Y79AA1002089	F-Y79AA1002089	0.94
	Y79AA1002093	F-Y79AA1002093	0.89
	Y79AA1002115	F-Y79AA1002115	0.31
25	Y79AA1002125	F-Y79AA1002125	0.76
	Y79AA1002139	F-Y79AA1002139	0. 32
	Y79AA1002204	F-Y79AA1002204	0. 94
30	Y79AA1002208	F-Y79AA1002208	0. 76
30	Y79AA1002209	F-Y79AA1002209	0.91
	Y79AA1002210	F-Y79AA1002210	0.60
	Y79AA1002229	F-Y79AA1002229	0.76
35	Y79AA1002234	F-Y79AA1002234	0.94
	Y79AA1002246	F-Y79AA1002246	0. 65
	Y79AA1002258	F-Y79AA1002258	0. 79
	Y79AA1002298	F-Y79AA1002298	0.83
40	Y79AA1002307	F-Y79AA1002307	0. 52
	Y79AA1002311	F-Y79AA1002311	0. 92
	Y79AA1002351	F-Y79AA1002351	0.85
	Y79AA1002361	F-Y79AA1002361	0. 90
45	Y79AA1002399	F-Y79AA1002399	0.94
	Y79AA1002407	F-Y79AA1002407	0.83
	Y79AA1002416	F-Y79AA1002416	0.77
50	Y79AA1002431	F-Y79AA1002431	0. 61
50	Y79AA1002433	F-Y79AA1002433	0.81
	Y79AA1002472	F-Y79AA1002472	0.39
	Y79AA1002482	F-Y79AA1002482	0.88
55	Y79AA1002487	F-Y79AA1002487	0.87

#### Table 10

Selected clones having the maximal ATGpr1 score of not more than 0.3 (1857 clones including the 9 clones whose 5'-end sequence does not contain the ATG codon. Blank of the maximal ATGpr1 score indicates that clones whose 5'-end sequence does not contain the ATG codon).

	clone	name of	maximal
	name :	sequence	ATGpr1
5	•		score
			····
	HEMBA100004	2 F-HEMBA1000	042 0.08
	HEMBA100011	1 F-HEMBA1000	111 0.1
10	HEMBA1000186	O F-HEMBA1000	180 0.09
	HEMBA100021	3 F-HEMBA1000	213 0. 21
	HEMBA100024	3 F-HEMBA1000	243 0.1
	HEMBA100025	1 F-HEMBA1000	251 0.08
15	HEMBA1000264	4 F-HEMBA1000	264 0.08
	HEMBA1000280	O F-HEMBA1000	280 0.13
	HEMBA1000282	2 F-HEMBA1000	282, 0.1
	HEMBA1000333		333 0.17
20	HEMBA1000338	8 F-HEMBA1000	338 0.16
	HEMBA100035	1 F-HEMBA1000	351 0.17
	HEMBA100035	5 F-HEMBA1000	
	HEMBA1000357		
25	HEMBA1000366	6 F-HEMBA1000	
	HEMBA1000376		
	HEMBA1000390		
	HEMBA1000411		111 0.07
30	HEMBA1000418	B F-HEMBA10004	
	HEMBA1000422		
	HEMBA1000428		
	HEMBA1000434		
25	HEMBA1000442		
35	HEMBA1000459		
	HEMBA1000464		
	HEMBA1000469		
	HEMBA1000504		• •
40	HEMBA1000518		
	HEMBA1000519		
	HEMBA1000540	_	
	HEMBA1000545		
45	HEMBA1000557		
	HEMBA1000563		
	HEMBA1000569		
	HEMBA1000575		
50	HEMBA1000604		
	HEMBA1000622		
	HEMBA1000655		
	HEMBA1000662		
55	HEMBA1000673		
	HEMBA1000702	2 F-HEMBA10007	02 0.12

	HEMBA1000705	F-HEMBA1000705	0. 06
	HEMBA1000722	F-HEMBA1000722	0. 17
	HEMBA1000726	F-HEMBA1000726	0. 08
5	HEMBA1000747	F-HEMBA1000747	0.06
	HEMBA1000749	F-HEMBA1000749	0.09
	HEMBA1000769	F-HEMBA1000769	0.19
	HEMBA1000773	F-HEMBA1000773	0. 08
10	HEMBA1000774	F-HEMBA1000774	0. 17
	HEMBA1000791	F-HEMBA1000791	0.08
	HEMBA1000822	F-HEMBA1000822	0. 19
	HEMBA1000827	F-HEMBA1000827	0.09
15	HEMBA1000843	F-HEMBA1000843	0. 12
	HEMBA1000852	F-HEMBA1000852	0.16
	HEMBA1000870	F-HEMBA1000870	0.14
	HEMBA1000876	F-HEMBA1000876	0. 15
20	HEMBA1000908	F-HEMBA1000908	0. 17
20	HEMBA1000934	F-HEMBA1000934	0. 2
	HEMBA1000943	F-HEMBA1000943	0.08
	HEMBA1000960	F-HEMBA1000960	0.09
	HEMBA1000972	F-HEMBA1000972	0. 12
25	HEMBA1000974	F-HEMBA1000974	0. 08
	HEMBA1000985	F-HEMBA1000985	0.06
	HEMBA1000991	F-HEMBA1000991	0. 23
	HEMBA1001007	F-HEMBA1001007	0. 06
30	HEMBA1001017	F-HEMBA1001017	0.06
	HEMBA1001019	F-HEMBA1001019	0. 08
	HEMBA1001020	F-HEMBA1001020	0. 15
	HEMBA1001024	F-HEMBA1001024	0. 1
35	HEMBA1001026	F-HEMBA1001026	0. 13
	HEMBA1001051	F-HEMBA1001051	0. 27
	HEMBA1001060	F-HEMBA1001060	0.11
	HEMBA1001071	F-HEMBA1001071	0. 1
40	HEMBA1001077	F-HEMBA1001077	0. 07
	HEMBA1001094	F-HEMBA1001094	0. 07
	HEMBA1001099	F-HEMBA1001099	0. 08
	HEMBA1001121	F-HEMBA1001121	0. 25
45	HEMBA1001123	F-HEMBA1001123	0. 05
	HEMBA1001172	F-HEMBA1001172	0. 08
	HEMBA1001208	F-HEMBA1001208	0. 11
	HEMBA1001226	F-HEMBA1001226	0. 27
50	HEMBA1001265	F-HEMBA1001265	0.06
50	HEMBA1001294	F-HEMBA1001294	0. 12
	HEMBA1001299	F-HEMBA1001299	0. 2
	HEMBA1001233	F-HEMBA1001319	0. 3
	HEMBA1001323	F-HEMBA1001323	0. 07
55	HEMBA1001327	F-HEMBA1001327	0. 08
	ILIND/(100102)	, ,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	J. VJ

HEMBA1001330 F-HEMBA1001330 HEMBA1001375 F-HEMBA1001375	
TEMBA10013/5 F-HEMBA10013/5	0.11
HENDALOOLOO E HENDALOO	0. 15
HEMBA1001383 F-HEMBA1001383	0. 11
UEMBY 1001331	<b></b> .
HEMBA1001411 F-HEMBA1001411	0.06
HEMBA1001432 F-HEMBA1001432	0. 07
HEMBA1001433 F-HEMBA1001433	0. 24
10 HEMBA1001435 F-HEMBA1001435	0.16
HEMBA1001442 F-HEMBA1001442	0. 22
HEMBA1001454 F-HEMBA1001454	0.11
HEMBA1001463 F-HEMBA1001463	0.08
15 HEMBA1001478 F-HEMBA1001478	0.13
HEMBA1001515 F-HEMBA1001515	0.3
HEMBA1001517 F-HEMBA1001517,	0.07
HEMBA1001522 F-HEMBA1001522	0.3
20 HEMBA1001557 F-HEMBA1001557	0.06
HEMBA1001566 F-HEMBA1001566	0.1
HEMBA1001585 F-HEMBA1001585	0.09
HEMBA1001589 F-HEMBA1001589	0.06
HFMRA1001608 F-HFMRA1001608	0. 17
HEMBA1001636 F-HEMBA1001636	0. 15
HEMBA1001651 F-HEMBA1001651	0. 23
HEMBA1001658 F-HEMBA1001658	0. 17
HEMBA1001675 F-HEMBA1001675	0. 11
30 HEMBA1001678 F-HEMBA1001678	0. 06
HEMBA1001681 F-HEMBA1001681	0.1
HEMBA1001709 F-HEMBA1001709	0. 29
HEMBA1001712 F-HEMBA1001712	0. 08
35 HEMBA1001718 F-HEMBA1001718	0. 07
HEMBA1001734 F-HEMBA1001734	0. 07
HEMBA1001745 F-HEMBA1001745	0. 09
HEMBA1001761 F-HEMBA1001761	0. 09
40 HEMBA1001784 F-HEMBA1001784	0. 11
HEMBA1001791 F-HEMBA1001791	0. 15
HEMBA1001803 F-HEMBA1001803	0. 13
HEMBA1001808 F-HEMBA1001808	0. 18
HEMRATOOT POO E-HEMRATOOT POO	0. 18
HEMBA1001835 F-HEMBA1001835	
	0. 29
	0.11
	0. 2
50	0. 18
MEMBERITATIO ( _ULBINATAMANA	0.09
HEMBA1001918 F-HEMBA1001918	41 73
HEMBA1001940 F-HEMBA1001940	0. 21
HEMBA1001940 F-HEMBA1001940 HEMBA1001942 F-HEMBA1001942	0.17
HEMBA1001940 F-HEMBA1001940	

	HEMBA1001962	F-HEMBA1001962	0.08
	HEMBA1001964	F-HEMBA1001964	0.15
	HEMBA1001979	F-HEMBA1001979	0.07
5	HEMBA1001991	F-HEMBA1001991	0.08
	HEMBA1002003	F-HEMBA1002003	0.11
	HEMBA1002008	F-HEMBA1002008	0.08
	HEMBA1002022	F-HEMBA1002022	0.08
10	HEMBA1002039	F-HEMBA1002039	0. 2
	HEMBA1002100	F-HEMBA1002100	0.07
	HEMBA1002113	F-HEMBA1002113	0.09
	HEMBA1002139	F-HEMBA1002139	0. 2
15	HEMBA1002144	F-HEMBA1002144	0.11
73	HEMBA1002153	F-HEMBA1002153	0.09
	HEMBA1002160	F-HEMBA1002160,	0, 15
	HEMBA1002162	F-HEMBA1002162	0. 19
	HEMBA1002166	F-HEMBA1002166	0.06
20	HEMBA1002185	F-HEMBA1002185	0. 11
	HEMBA1002204	F-HEMBA1002204	0.09
	HEMBA1002226	F-HEMBA1002226	0.13
	HEMBA1002253	F-HEMBA1002253	0.05
25	HEMBA1002257	F-HEMBA1002257	0.09
	HEMBA1002270	F-HEMBA1002270	0.06
	HEMBA1002321	F-HEMBA1002321	0. 07
	HEMBA1002328	F-HEMBA1002328	0.06
30	HEMBA1002337	F-HEMBA1002337	0.11
	HEMBA1002348	F-HEMBA1002348	0. 12
	HEMBA1002349	F-HEMBA1002349	0. 05
	HEMBA1002381	F-HEMBA1002381	0.09
35	HEMBA1002389	F-HEMBA1002389	0. 14
	HEMBA1002486	F-HEMBA1002486	0. 1
	HEMBA1002498	F-HEMBA1002498	0.17
	HEMBA1002538	F-HEMBA1002538	0. 27
40	HEMBA1002552	F-HEMBA1002552	0. 23
	HEMBA1002555	F-HEMBA1002555	0. 11
	HEMBA1002558	F-HEMBA1002558	0. 11
	HEMBA1002561	F-HEMBA1002561	0. 11
45	HEMBA1002590	F-HEMBA1002590	0.06
	HEMBA1002592	F-HEMBA1002592	0. 17
	HEMBA1002621	F-HEMBA1002621	0. 12
	HEMBA1002628	F-HEMBA1002628	0.06
50	HEMBA1002629 HEMBA1002645	F-HEMBA1002629	0. 23
		F-HEMBA1002645	0. 12
	HEMBA1002651 HEMBA1002659	F-HEMBA1002651	0.08
	HEMBA1002661	F-HEMBA1002659 F-HEMBA1002661	0. 28 0. 28
55	HEMBA1002666	F-HEMBA1002666	
JJ	IILHUM I UUZOOD	1 TIEMDA 1002000	0. 16

	HEMBA1002678	F-HEMBA1002678	0. 07
	HEMBA1002679	F-HEMBA1002679	0. 14
_	HEMBA1002712	F-HEMBA1002712	0. 26
5	HEMBA1002716	F-HEMBA1002716	_0.14
	HEMBA1002728	F-HEMBA1002728	0.09
	HEMBA1002730	F-HEMBA1002730	0. 07
	HEMBA1002742	F-HEMBA1002742	0.09
10	HEMBA1002748	F-HEMBA1002748	0. 17
	HEMBA1002780	F-HEMBA1002780	0. 1
	HEMBA1002801	F-HEMBA1002801	0. 1
	HEMBA1002826	F-HEMBA1002826	0. 1
15	HEMBA1002833	F-HEMBA1002833	0.14
	HEMBA1002886	F-HEMBA1002886	0.09
	HEMBA1002896	F-HEMBA1002896	
	HEMBA1002921	F-HEMBA1002921	0. 25
20	HEMBA1002924	F-HEMBA1002924	
	HEMBA1002934	F-HEMBA1002934	
	HEMBA1002944		
	HEMBA1002968		0. 12
25	HEMBA1003034		0.06
·	HEMBA1003037		0.11
	HEMBA1003064 HEMBA1003071		0.09
	HEMBA1003071	F-HEMBA1003071 F-HEMBA1003083	0. 27 0. 18
30	HEMBA1003086	F-HEMBA1003086	0. 16
	HEMBA1003098	F-HEMBA1003098	0. 00
	HEMBA1003133	F-HEMBA1003133	0. 12
	HEMBA1003142	F-HEMBA1003142	0. 29
35	HEMBA1003166	F-HEMBA1003166	0. 1
	HEMBA1003197	F-HEMBA1003197	0. 1
	HEMBA1003202	F-HEMBA1003202	0. 17
	HEMBA1003204	F-HEMBA1003204	0. 13
40	HEMBA1003212	F-HEMBA1003212	0.13
	HEMBA1003220	F-HEMBA1003220	0.14
	HEMBA1003229	F-HEMBA1003229	0. 19
	HEMBA1003273	F-HEMBA1003273	0. 1
45	HEMBA1003276	F-HEMBA1003276	0. 12
	HEMBA1003278	F-HEMBA1003278	0.09
	HEMBA1003296	F-HEMBA1003296	0. 07
	HEMBA1003304	F-HEMBA1003304	0.09
50	HEMBA1003309	F-HEMBA1003309	0. 07
	HEMBA1003314	F-HEMBA1003314	0. 21
	HEMBA1003328	F-HEMBA1003328	0. 13
	HEMBA1003330	F-HEMBA1003330	0. 08
55	HEMBA1003348	F-HEMBA1003348	0. 17
•	HEMBA1003373	F-HEMBA1003373	0. 21

	HEMBA1003376 F-HEMBA1003376 0	. 14
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		. 08
30		. 08
		. 09
		. 2
		. 12
35	HEMBA1003942 F-HEMBA1003942 0	. 1
	HEMBA1003950 F-HEMBA1003950 0	. 09
	HEMBA1003958 F-HEMBA1003958 0	. 07
	HEMBA1003976 F-HEMBA1003976 0	. 28
40	HEMBA1003987 F-HEMBA1003987 0	. 24
	HEMBA1004000 F-HEMBA1004000 0	. 16
	HEMBA1004012 F-HEMBA1004012 0	. 12
	HEMBA1004015 F-HEMBA1004015 0	. 23
45	HEMBA1004024 F-HEMBA1004024 0	. 13
		. 09
		. 09
		. 09
50		. 09
-		. 08
		. 07
		. 08
55		. 1
JJ	HEMBA1004241 F-HEMBA1004241 0	. 08

	HEMBA1004267	F-HEMBA1004267	0.09
	HEMBA1004272	F-HEMBA1004272	0. 09
	HEMBA1004295	F-HEMBA1004295	0. 19
5	HEMBA1004306	F-HEMBA1004306	
	HEMBA1004312	F-HEMBA1004312	0. 26
	HEMBA1004323	F-HEMBA1004323	0. 13
	HEMBA1004334	F-HEMBA1004334	0.06
10	HEMBA1004354	F-HEMBA1004354	0. 1
	HEMBA1004356	F-HEMBA1004356	0.07
	HEMBA1004366	F-HEMBA1004366	0.14
	HEMBA1004394	F-HEMBA1004394	0.08
15	HEMBA1004396	F-HEMBA1004396	0.05
75	HEMBA1004405	F-HEMBA1004405	0. 17
	HEMBA1004429	F-HEMBA1004429,	
	HEMBA1004433	F-HEMBA1004433	0. 29
	HEMBA1004460	F-HEMBA1004460	0.11
20	HEMBA1004461	F-HEMBA1004461	0.11
	HEMBA1004482	F-HEMBA1004482	0. 15
	HEMBA1004506	F-HEMBA1004506	0.05
	HEMBA1004538	F-HEMBA1004538	0. 15
25	HEMBA1004577	F-HEMBA1004577	0.17
	HEMBA1004586	F-HEMBA1004586	0.06
	HEMBA1004617	F-HEMBA1004617	0.11
	HEMBA1004629	F-HEMBA1004629	0:11
30	HEMBA1004631	F-HEMBA1004631	0.1
	HEMBA1004666	F-HEMBA1004666	0.08
	HEMBA1004670	F-HEMBA1004670	0.15
	HEMBA1004672	F-HEMBA1004672	0.14
35	HEMBA1004730	F-HEMBA1004730	0. 15
	HEMBA1004733	F-HEMBA1004733	0.12
	HEMBA1004748	F-HEMBA1004748	0.1
	HEMBA1004770	F-HEMBA1004770	0.15
40	HEMBA1004778	F-HEMBA1004778	0.14
	HEMBA1004803	F-HEMBA1004803	0.13
	HEMBA1004807	F-HEMBA1004807	0.18
	HEMBA1004816	F-HEMBA1004816	0.06
45	HEMBA1004820	F-HEMBA1004820	0.11
	HEMBA1004865	F-HEMBA1004865	0. 17
	HEMBA1004880	F-HEMBA1004880	0. 28
	HEMBA1004900	F-HEMBA1004900	0.11
50	HEMBA1004909	F-HEMBA1004909	0.11
	HEMBA1004918	F-HEMBA1004918	0.09
	HEMBA1004956	F-HEMBA1004956	0.05
	HEMBA1004960	F-HEMBA1004960	0. 21
	HEMBA1004978	F-HEMBA1004978	0. 1
55	HEMBA1004983	F-HEMBA1004983	0. 28

	HEMBA1004995	F-HEMBA1004995	0. 24
	HEMBA1005008	F-HEMBA1005008	0. 24
	HEMBA1005039	F-HEMBA1005039	0. 07
5	HEMBA1005062		0.08
	~	F-HEMBA1005062	
	HEMBA1005123	F-HEMBA1005123	0.07
	HEMBA1005152	F-HEMBA1005152	0.11
10	HEMBA1005159	F-HEMBA1005159	0.1
70	HEMBA1005232	F-HEMBA1005232	0. 1
	HEMBA1005241	F-HEMBA1005241	0. 14
	HEMBA1005251	F-HEMBA1005251	0. 29
	HEMBA1005274	F-HEMBA1005274	0.06
15	HEMBA1005275	F-HEMBA1005275	0. 12
	HEMBA1005293	F-HEMBA1005293	0.06
	HEMBA1005304	F-HEMBA1005304,	
	HEMBA1005311	F-HEMBA1005311	0. 2
20	HEMBA1005315	F-HEMBA1005315	0.07
	- HEMBA1005318	F-HEMBA1005318	0.08
	HEMBA1005353	F-HEMBA1005353	0. 12
	HEMBA1005359	F-HEMBA1005359	0. 15
25	HEMBA1005372	F-HEMBA1005372	0. 07
	HEMBA1005374	F-HEMBA1005374	0.09
	HEMBA1005389	F-HEMBA1005389	0. 1
	HEMBA1005408	F-HEMBA1005408	0.09
30	HEMBA1005410	F-HEMBA1005410	0. 07
	HEMBA1005426	F-HEMBA1005426	0.16
	HEMBA1005443	F-HEMBA1005443	0. 17
	HEMBA1005447	F-HEMBA1005447	0. 25
	HEMBA1005497	F-HEMBA1005497	0.09
35	HEMBA1005500	F-HEMBA1005500	0.08
	HEMBA1005506	F-HEMBA1005506	0.17
	HEMBA1005508	F-HEMBA1005508	0. 12
	HEMBA1005511	F-HEMBA1005511	0.1
40	HEMBA1005520	F-HEMBA1005520	0.07
	HEMBA1005552	F-HEMBA1005552	0. 05
	HEMBA1005568	F-HEMBA1005568	0.06
	HEMBA1005570	F-HEMBA1005570	0. 1
45	HEMBA1005577	F-HEMBA1005577	0. 13
	HEMBA1005588	F-HEMBA1005588	0. 25
	HEMBA1005593	F-HEMBA1005593	0.13
	HEMBA1005606	F-HEMBA1005606	0. 16
50	HEMBA1005616	F-HEMBA1005616	0. 15
	HEMBA1005627	F-HEMBA1005627	0. 11
	HEMBA1005631	F-HEMBA1005631	0. 07
	HEMBA1005632	F-HEMBA1005632	0. 07
55	HEMBA1005634	F-HEMBA1005634	0. 07
	HEMBA1005670	F-HEMBA1005670	0. 11

	HEMBA1005679 F-HEMBA1005679	0. 22
	HEMBA1005699 F-HEMBA1005699	0.08
	HEMBA1005705 F~HEMBA1005705	0.08
5	HEMBA1005717 F-HEMBA1005717	0.17
	HEMBA1005765 F-HEMBA1005765	0.08
	HEMBA1005780 F-HEMBA1005780	0.16
	HEMBA1005829 F-HEMBA1005829	0.09
10	HEMBA1005853 F-HEMBA1005853	0.06
	HEMBA1005894 F-HEMBA1005894	0.16
	HEMBA1005909 F-HEMBA1005909	0.13
	HEMBA1005911 F-HEMBA1005911	0. 22
15	HEMBA1005921 F-HEMBA1005921	0. 1
	HEMBA1005931 F-HEMBA1005931	0.16
	HEMBA1005934 F-HEMBA1005934,	
	HEMBA1005962 F-HEMBA1005962	0.07
20	HEMBA1005999 F-HEMBA1005999	0.14
20	HEMBA1006002 F-HEMBA1006002	0.09
	HEMBA1006005 F-HEMBA1006005	0. 2
	HEMBA1006036 F-HEMBA1006036	0. 24
	HEMBA1006042 F-HEMBA1006042	0.11
25	HEMBA1006067 F-HEMBA1006067	0.09
	HEMBA1006090 F-HEMBA1006090	0.09
	HEMBA1006124 F-HEMBA1006124	0.05
	HEMBA1006142 F-HEMBA1006142	0.13
30	HEMBA1006158 F-HEMBA1006158	0. 27
	HEMBA1006253 F-HEMBA1006253	0. 29
	HEMBA1006259 F-HEMBA1006259	0.11
	HEMBA1006268 F-HEMBA1006268	0.11
35	HEMBA1006328 F-HEMBA1006328	0.06
	HEMBA1006334 F-HEMBA1006334	0.08
	HEMBA1006359 F-HEMBA1006359	0.22
	HEMBA1006364 F-HEMBA1006364	0.23
40	HEMBA1006380 F-HEMBA1006380	0.09
	HEMBA1006416 F-HEMBA1006416	0. 25
	HEMBA1006419 F-HEMBA1006419	0.08
	HEMBA1006421 F-HEMBA1006421	0.26
45	HEMBA1006426 F-HEMBA1006426	0.17
	HEMBA1006438 F-HEMBA1006438	0.08
	HEMBA1006446 F-HEMBA1006446	0.11
	HEMBA1006461 F-HEMBA1006461	0.15
50	HEMBA1006471 F-HEMBA1006471	0.06
J.	HEMBA1006486 F-HEMBA1006486	0.11
	HEMBA1006489 F-HEMBA1006489	0.09
	HEMBA1006540 F-HEMBA1006540	0. 23
EF	HEMBA1006546 F-HEMBA1006546	0.07
55	HEMBA1006562 F-HEMBA1006562	0.12

	HEMBA1006569	F-HEMBA1006569	
	HEMBA1006595	F-HEMBA1006595	0. 08
	HEMBA1006597	F-HEMBA1006597	0.17
5	HEMBA1006617	F-HEMBA1006617	0.07
	HEMBA1006635	F-HEMBA1006635	0.07
	HEMBA1006639	F-HEMBA1006639	0. 25
	HEMBA1006648	F-HEMBA1006648	0. 24
10	HEMBA1006653	F-HEMBA1006653	0. 09
	HEMBA1006665	F-HEMBA1006665	0. 17
	HEMBA1006676	F-HEMBA1006676	0.17
	HEMBA1006695	F-HEMBA1006695	0. 1
15	HEMBA1006696	F-HEMBA1006696	0. 05
	HEMBA1006744	F-HEMBA1006744	0.12
	HEMBA1006779	F-HEMBA1006779	0.07
	HEMBA1006780	F-HEMBA1006780	0. 23
20	HEMBA1006821	F-HEMBA1006821	0.08
	HEMBA1006824	F-HEMBA1006824	0.09
	HEMBA1006849	F~HEMBA1006849	0.05
	HEMBA1006865	F-HEMBA1006865	0. 1
25	HEMBA1006921	F~HEMBA1006921	0. 22
	HEMBA1006929	F-HEMBA1006929	0. 15
	HEMBA1006938	F-HEMBA1006938	0. 07
	HEMBA1006949	F-HEMBA1006949	0. 27
30	HEMBA1007017	F-HEMBA1007017	0. 07
	HEMBA1007045	F-HEMBA1007045	0. 14
	HEMBA1007051 HEMBA1007073	F-HEMBA1007051	0. 19
	HEMBA1007078	F-HEMBA1007073 F-HEMBA1007078	0. 22 0. 12
35	HEMBA1007113	F-HEMBA1007113	0. 12
	HEMBA1007113	F-HEMBA1007129	0. 12
	HEMBA1007147	F-HEMBA1007147	0. 18
	HEMBA1007206	F-HEMBA1007206	0. 11
40	HEMBA1007256	F-HEMBA1007256	0. 13
	HEMBA1007273	F-HEMBA1007273	0. 05
	HEMBA1007279	F-HEMBA1007279	0.09
	HEMBA1007288	F-HEMBA1007288	0.06
45	HEMBA1007322	F-HEMBA1007322	0. 12
40	HEMBA1007327	F-HEMBA1007327	0.12
	HEMBA1007341	F-HEMBA1007341	0.1
	HEMBA1007347	F-HEMBA1007347	0.14
50	HEMBB1000005	F-HEMBB1000005	0. 13
50	HEMBB1000039	F-HEMBB1000039	0.07
	HEMBB1000044	F-HEMBB1000044	0.08
	HEMBB1000050	F-HEMBB1000050	0. 07
44	HEMBB1000054	F-HEMBB1000054	0. 12
55	HEMBB1000055	F-HEMB81000055	0. 2

	HEMBB1000059	E-UENDD100050	0.00
	• • • • • • • • • • • • • • • • • • • •	F-HEMBB1000059	0.08
	HEMBB1000089	F-HEMBB1000089	0.09
5	HEMBB1000099	F-HEMBB1000099	0.17
3	HEMBB1000113	F-HEMBB1000113	0. 12
	HEMBB1000141	F-HEMBB1000141	0. 13
	HEMBB1000144	F-HEMBB1000144	0. 2
	HEMBB1000218	F-HEMBB1000218	0. 07
10	HEMBB1000250	F-HEMBB1000250	0. 26
	HEMBB1000258	F-HEMBB1000258	0. 15
	HEMBB1000264	F-HEMBB1000264	0.08
	HEMB81000272	F-HEMBB1000272	0.11
15	HEMBB1000274	F-HEMBB1000274	0. 12
	HEMBB1000284	F-HEMBB1000284	0. 1
	HEMBB1000307	F-HEMBB1000307,	
	HEMBB1000312	F-HEMBB1000312	0.07
20	HEMBB1000317	F-HEMBB1000317	0. 27
	HEMBB1000318	F-HEMBB1000318	0.11
	HEMBB1000335	F-HEMBB1000335	0.15
	HEMBB1000336	F-HEMBB1000336	0. 28
25	HEMBB1000337	F-HEMBB1000337	0. 25
20	HEMBB1000341	F-HEMBB1000341	0. 24
	HEMBB1000343	F-HEMBB1000343	0.07
	HEMBB1000354	F-HEMBB1000354	0. 15
	HEMBB1000369	F-HEMBB1000369	0.08
30	HEMBB1000374	F-HEMBB1000374	0.3
·	HEMBB1000376	F-HEMBB1000376	0. 12
	HEMBB1000399	F-HEMBB1000399	0.08
	HEMBB1000402	F-HEMBB1000402	0. 29
35	HEMBB1000404	F-HEMBB1000404	0. 29
	HEMBB1000420	F-HEMBB1000420	0.11
	HEMBB1000434	F-HEMBB1000434	0.05
	HEMBB1000441	F-HEMBB1000441	0. 12
40	HEMBB1000455	F-HEMBB1000455	0.05
	HEMBB1000472		0.05
	HEMBB1000480		0.13
	HEMBB1000487	F-HEMBB1000487	0.09
45	HEMBB1000490	F-HEMBB1000490	0.06
	HEMBB1000491	F-HEMBB1000491	0.16
	HEMBB1000493	F-HEMBB1000493	0. 15
	HEMBB1000518	F-HEMBB1000518	0.05
50	HEMBB1000523	F-HEMBB1000523	0.07
	HEMBB1000530	F-HEMBB1000530	0.14
	HEMBB1000554	F-HEMBB1000554	0. 23
	HEMBB1000564	F-HEMBB1000564	0.09
	HEMBB1000573	F-HEMBB1000573	0.1
55	HEMBB1000575	F-HEMBB1000575	0.08

	HEMBB1000586	F-HEMBB1000586	0.·12
	HEMBB1000598	F-HEMBB1000598	0.12
	HEMBB1000637	F-HEMBB1000637	0.19
5	HEMBB1000638	F-HEMBB1000638	0.07
	HEMBB1000643	F-HEMBB1000643	0.07
	HEMBB1000649	F-HEMBB1000649	0.13
	HEMBB1000652	F-HEMBB1000652	0. 2
10	HEMBB1000665	F-HEMBB1000665	0.07
	HEMBB1000684	F-HEMBB1000684	0.06
	HEMBB1000709	F-HEMBB1000709	0.09
	HEMBB1000726	F-HEMBB1000726	0.07
15	HEMBB1000738	F-HEMBB1000738	0. 19
73	HEMBB1000749	F-HEMBB1000749	0. 14
	HEMBB1000770	F-HEMBB1000770,	
	HEMBB1000790	F-HEMBB1000790	0. 23
	HEMBB1000794	F-HEMBB1000794	0.09
20	HEMBB1000821	F-HEMBB1000821	0.1
	HEMBB1000822	F-HEMBB1000822	0.18
	HEMBB1000827	F-HEMBB1000827	0.1
•	HEMBB1000831	F-HEMBB1000831	0. 2
25	HEMBB1000840	F-HEMBB1000840	0.09
	HEMBB1000876	F-HEMBB1000876	0.12
	HEMBB1000883	F-HEMB81000883	0.13
	HEMBB1000888	F-HEMBB1000888	0.1
30	HEMBB1000890	F-HEMBB1000890	0.09
	HEMBB1000893	F-HEMBB1000893	0.1
	HEMBB1000910	F-HEMBB1000910	0.06
	HEMBB1000913	F-HEMBB1000913	0.09
35	HEMB81000915	F-HEMBB1000915	0.12
	HEMBB1000917	F-HEMBB1000917	0.15
	HEMBB1000959	F-HEMBB1000959	0.08
	HEMBB1000981	F-HEMBB1000981	0.09
40	HEMBB1000996	F-HEMBB1000996	0.1
	HEMBB1001004	F-HEMB81001004	0.08
	HEMBB1001008	F-HEMBB1001008	0. 25
	HEMBB1001037	F-HEMBB1001037	0.14
45	HEMBB1001047	F-HEMBB1001047	0.14
45	HEMBB1001051	F-HEMBB1001051	0.13
	HEMBB1001060	F-HEMBB1001060	0.16
	HEMBB1001063	F-HEMBB1001063	0.1
	HEMBB1001102	F-HEMBB1001102	0.19
50	HEMBB1001114	F-HEMBB1001114	0.08
	HEMBB1001119	F-HEMBB1001119	0.13
	HEMBB1001133	F-HEMBB1001133	0.06
	HEMBB1001142	F-HEMBB1001142	0.17
55	HEMB81001177	F-HEMBB1001177	0. 2

	HEMBB1001208	F-HEMBB1001208	0.12
	HEMBB1001209	F-HEMBB1001209	0.11
	HEMBB1001218	F-HEMBB1001218	0.1
5	HEMBB1001221	F-HEMBB1001221	0.1
	HEMBB1001249	F-HEMBB1001249	0. 12
	HEMBB1001253	F-HEMBB1001253	0.13
	HEMBB1001254	F-HEMB81001254	0.13
10	HEMBB1001267	F-HEMBB1001267	0.09
	HEMBB1001271	F-HEMBB1001271	0.16
	HEMBB1001282	F-HEMBB1001282	0.14
	HEMBB1001302	F-HEMBB1001302	0.18
15	HEMBB1001304	F-HEMBB1001304	0.17
	HEMBB1001315	F-HEMBB1001315	0.18
	HEMBB1001317	F-HEMBB1001317,	0.12
	HEMBB1001326	F-HEMBB1001326	0.09
20	HEMBB1001335	F-HEMBB1001335	0.08
20	HEMBB1001337	F-HEMBB1001337	0.12
	HEMBB1001348	F-HEMBB1001348	0.11
	HEMBB1001356	F-HEMBB1001356	0.1
05	HEMBB1001366	F-HEMBB1001366	0.16
25	HEMBB1001367	F-HEMBB1001367	0.11
	HEMBB1001380	F-HEMBB1001380	0.07
	HEMBB1001424	F-HEMBB1001424	0.05
	HEMBB1001436	F-HEMBB1001436	0.14
30	HEMBB1001443	F-HEMBB1001443	0. 28
	HEMBB1001454	F-HEMBB1001454	
	HEMBB1001458	F-HEMBB1001458	0.11
	HEMBB1001463	F-HEMBB1001463	0.1
35	HEMBB1001464	F-HEMBB1001464	0.06
	HEMBB1001500	F-HEMBB1001500	0.11
	HEMBB1001521	F-HEMB81001521	0.13
	HEMBB1001527	F-HEMBB1001527	0.13
40	HEMBB1001535	F-HEMBB1001535	0.11
	HEMBB1001536	F-HEMBB1001536	0.08
	HEMBB1001537	F-HEMBB1001537	0.11
	HEMBB1001555	F-HEMBB1001555	0.08
45	HEMBB1001565	F-HEMBB1001565	0.1
	HEMBB1001586	F-HEMBB1001586	0.06
	HEMBB1001588	F-HEMB81001588	0.06
	HEMBB1001618	F-HEMBB1001618	0.09
50	HEMBB1001619	F-HEMBB1001619	0.12
	HEMBB1001630	F-HEMBB1001630	0.12
	HEMBB1001637	F-HEMBB1001637	0.08
	HEMBB1001641	F-HEMBB1001641	0.09
55	HEMBB1001704	F-HEMBB1001704	0.09
	HEMBB1001706	F-HEMBB1001706	0. 11

	HEMBB1001717	F-HEMBB1001717	0.09
	HEMBB1001735	F-HEMBB1001735	0. 21
	HEMBB1001747	F-HEMBB1001747	0.07
5	HEMBB1001749	F-HEMBB1001749	0, 13
	HEMBB1001753	F-HEMBB1001753	0.08
	HEMB81001756	F-HEMBB1001756	0.11
	HEMBB1001762	F-HEMBB1001762	0.11
10	HEMBB1001797	F-HEMBB1001797	0.13
	HEMBB1001802	F-HEMBB1001802	0.06
	HEMBB1001836	F-HEMBB1001836	0.15
	HEMB81001850	F-HEMBB1001850	0.06
15	HEMBB1001863	F-HEMBB1001863	0.09
	HEMBB1001867	F-HEMBB1001867	0.11
	HEMBB1001868	F-HEMBB1001868,	0. 24
	HEMBB1001875	F-HEMBB1001875	0.05
20	HEMBB1001880	F-HEMBB1001880	0.08
20	HEMBB1001899	F-HEMBB1001899	0.16
	HEMBB1001911	F-HEMBB1001911	0.22
	HEMBB1001921	F-HEMBB1001921	0.07
05	HEMBB1001922	F-HEMBB1001922	0.11
25	HEMBB1001925	F-HEMBB1001925	0.19
	HEMB81001930	F-HEMBB1001930	0.1
	HEMBB1001944	F-HEMBB1001944	0.14
	HEMBB1001952	F-HEMBB1001952	0.08
30	HEMBB1001953	F-HEMBB1001953	0.15
	HEMBB1001967	F-HEMBB1001967	0.08
	HEMBB1001973	F-HEMBB1001973	0.09
	HEMBB1001983	F-HEMBB1001983	0. 1
35	HEMBB1001988	F-HEMBB1001988	0.15
	HEMBB1001996	F-HEMBB1001996	0.19
	HEMBB1001997	F-HEMBB1001997	0.08
	HEMBB1002002	F-HEMBB1002002	0. 21
40	HEMBB1002005	F-HEMBB1002005	0.18
	HEMBB1002009	F-HEMBB1002009	0.09
	HEMBB1002015		0. 26
	HEMBB1002042	F-HEMBB1002042	0.11
45	HEMBB1002043	F-HEMBB1002043	0.09
	HEMBB1002045	F-HEMBB1002045	0. 26
	HEMBB1002049	F-HEMBB1002049	0.12
	HEMBB1002069	F-HEMBB1002069	0.12
50	HEMBB1002092	F-HEMBB1002092	0.16
	HEMBB1002094	F-HEMBB1002094	0.11
	HEMBB1002115	F-HEMBB1002115	0.09
	HEMBB1002139	F-HEMB81002139	0.13
EE	HEMBB1002189	F-HEMBB1002189	0.06
55	HEMBB1002190	F-HEMBB1002190	0.18

	HEMBB1002232	F-HEMBB1002232	0.09
	HEMBB1002247	F-HEMBB1002247	0.17
	HEMBB1002254	F-HEMBB1002254	0.05
5	HEMBB1002255	F-HEMBB1002255	0.11
	HEMBB1002266	F-HEMBB1002266	0. 28
	HEMBB1002280	F-HEMBB1002280	0.07
	HEMBB1002306	F-HEMBB1002306	0.06
10	HEMBB1002364	F-HEMBB1002364	0.05
	HEMBB1002371	F-HEMBB1002371	0.12
	HEMBB1002381	F-HEMBB1002381	0.09
	HEMBB1002387	F-HEMBB1002387	0.17
15	HEMBB1002425	F-HEMBB1002425	0.08
·-	HEMBB1002453	F-HEMBB1002453	0.12
	HEMBB1002458	F-HEMBB1002458,	
	HEMBB1002477	F-HEMBB1002477	0. 17
22	HEMBB1002509	F-HEMBB1002509	0.06
20	HEMBB1002520	F-HEMBB1002520	0.08
	HEMBB1002522	F-HEMBB1002522	0. 21
	HEMBB1002531	F-HEMBB1002531	0. 1
	HEMBB1002534	F-HEMBB1002534	0.09
25	HEMBB1002556	F-HEMBB1002556	0.1
	HEMBB1002579	F-HEMBB1002579	0. 13
	HEMBB1002582	F-HEMBB1002582	0. 17
	HEMBB1002590	F-HEMBB1002590	0.08
30	HEMBB1002596	F-HEMBB1002596	0. 15
	HEMBB1002601	F-HEMBB1002601	0.1
	HEMBB1002603	F-HEMBB1002603	0. 12
	HEMBB1002610	F-HEMBB1002610	0. 1
35	HEMBB1002613	F-HEMBB1002613	0.09
	HEMBB1002617	F-HEMBB1002617	0. 12
	HEMBB1002623	F-HEMBB1002623	0.14
	HEMBB1002635	F-HEMBB1002635	0.08
40	HEMBB1002664	F-HEMBB1002664	0.06
	HEMBB1002677	F-HEMBB1002677	0.07
	HEMBB1002683	F-HEMBB1002683	0.06
	HEMBB1002686	F-HEMBB1002686	0.12
45	HEMBB1002699	F-HEMBB1002699	0.15
45	HEMBB1002702	F-HEMBB1002702	0.16
	HEMBB1002712	F-HEMBB1002712	0.07
	MAMMA1000009	F-MAMMA1000009	0.11
	MAMMA1000043	F-MAMMA1000043	0.19
50	MAMMA1000045	F-MAMMA1000045	0.08
	MAMMA1000092	F-MAMMA1000092	0.11
	MAMMA1000103	F-MAMMA1000103	0.09
	MAMMA1000117	F-MAMMA1000117	0. 2
55	MAMMA1000129	F-MAMMA1000129	0. 17
			•

	MAMMA1000133	F-MAMMA1000133	0. 1
	MAMMA1000134	F-MAMMA1000134	0. 05
	MAMMA1000155	F-MAMMA1000155	0. 12
5	MAMMA1000175	F-MAMMA1000175	0. 07
	MAMMA1000198	F-MAMMA1000198	0. 2
	MAMMA1000221	F-MAMMA1000221	0. 1
	MAMMA1000227	F-MAMMA1000227	0. 08
10	MAMMA1000241	F-MAMMA1000241	0. 16
	MAMMA1000254	F-MAMMA1000254	0.08
	MAMMA1000264	F-MAMMA1000264	0. 1
	MAMMA1000266	F-MAMMA1000266	0.18
45	MAMMA1000270	F-MAMMA1000270	0. 1
15	MAMMA1000273	F-MAMMA1000287	0. 1
	MAMMA1000302	F-MAMMA1000302,	
	MAMMA1000307	F-MAMMA1000307	0. 26
	MAMMA1000331	F-MAMMA1000331	0. 18
20	MAMMA1000340	F-MAMMA1000340	0. 2
	MAMMA1000348	F-MAMMA1000348	0. 09
	MAMMA1000356	F-MAMMA1000356	0. 15
	MAMMA1000360	F-MAMMA1000360	0.09
25	MAMMA1000385	F-MAMMA1000385	0. 08
	MAMMA1000402	F-MAMMA1000402	0.11
	MAMMA1000413	F-MAMMA1000413	0.1
	MAMMA1000414	F-MAMMA1000414	0.1
30	MAMMA1000423	F-MAMMA1000423	0.06
	MAMMA1000424	F-MAMMA1000424	0. 1
	MAMMA1000431	F-MAMMA1000431	0. 28
	MAMMA1000444	F-MAMMA1000444	0.07
35	MAMMA1000446	F-MAMMA1000446	0.08
	MAMMA1000478	F-MAMMA1000478	0. 1
	MAMMA1000483	F-MAMMA1000483	0.07
	MAMMA1000500	F-MAMMA1000500	0.16
40	MAMMA1000501	F-MAMMA1000501	0.05
	MAMMA1000516	F-MAMMA1000516	0.06
	MAMMA1000522	F-MAMMA1000522	0. 21
	MAMMA1000559	F-MAMMA1000559	0.12
45	MAMMA1000565	F-MAMMA1000565	0.09
-	MAMMA1000576	F-MAMMA1000576	0.14
	MAMMA1000585	F-MAMMA1000585	0. 07
	MAMMA1000594	F-MAMMA1000594	0. 17
50	MAMMA1000597	F-MAMMA1000597	0.1
50	MAMMA1000605	F-MANIMA1000605	0.09
	MAMMA1000616	F-MAMMA1000616	0.08
	MAMMA1000621	F-MAMMA1000621	0. 07
	MAMMA1000643	F-MAMMA1000643	0. 1
55	MAMMA1000669	F-MAMMA1000669	0. 11

	MAMMA1000696	F-MAMMA1000696	0.08
	MAMMA1000707	F-MAMMA1000707	0.07
	MAMMA1000714	F-MAMMA1000714	0.18
5	MAMMA1000718	F-MAMMA1000718	0.06
	MAMMA1000720	F-MAMMA1000720	0. 15
	MAMMA1000723	F-MAMMA1000723	0.14
	MAMMA1000732	F-MAMMA1000732	0.06
10	MAMMA1000733	F-MAMMA1000733	0. 1
	MAMMA1000744	F-MAMMA1000744	0. 15
	MAMMA1000752	F-MAMMA1000752	0.12
	MAMMA1000761	F-MAMMA1000761	0. 24
15	MAMMA1000775	F-MAMMA1000775	0, 24
	MAMMA1000778	F-MAMMA1000778	0. 09
	MAMMA1000782	F-MAMMA1000782,	0. 1
	MAMMA1000798	F-MAMMA1000798	0.16
20	MAMMA1000802	F-MAMMA1000802	0.11
20	MAMMA1000839	F-MAMMA1000839	0.09
	MAMMA1000845	F-MAMMA1000845	0.1
	MAMMA1000851	F-MAMMA1000851	0.08
	MAMMA1000855	F-MAMMA1000855	0. 23
25	MAMMA1000862	F-MAMMA1000862	0.12
	MAMMA1000863	F-MAMMA1000863	0.16
	MAMMA1000867	F-MAMMA1000867	0. 25
	MAMMA1000876	F-MAMMA1000876	0.14
30	MAMMA1000877	F-MAMMA1000877	0.07
	MAMMA1000880	F-MAMMA1000880	0. 13
	MAMMA1000883	F-MAMMA1000883	0.3
	MAMMA1000905	F-MANMA1000905	0.08
35	MAMMA1000931	F-MAMMA1000931	0.18
	MAMMA1000940	F-MAMMA1000940	0. 21
	MAMMA1000941	F-MAMMA1000941	0.13
	MAMMA1000942	F-MAMMA1000942	0.08
40	MAMMA1000943	F-MAMMA1000943	0.09
	MAMMA1000957	F-MAMMA1000957	0.13
	MAMMA1000962	F-MAMMA1000962	0.06
	MAMMA1000975	F-MAMMA1000975	0. 26
45	MAMMA1000987	F-MAMMA1000987	0.08
	MAMMA1000998	F-MAMMA1000998	0.08
	MAMMA1001003	F-MAMMA1001003	0. 15
	MAMMA1001024	F-MAMMA1001024	0. 08
50	MAMMA1001030	F-MAMMA1001030	0. 2
	MAMMA1001035	F-MAMMA1001035	0.08
	MAMMA1001038	F-MAMMA1001038	0. 15
	MAMMA1001050	F-MAMMA1001050	0.06
6.5	MAMMA1001067	F-MANMA1001067	0.08
55	MAMMA1001074	F-MAMMA1001074	0.09

	MAMMA1001082	F-MAMMA1001082	0.1
	MAMMA1001092	F-MAMMA1001092	0.05
	MAMMA1001133	F-MAMMA1001133	0.11
5	MAMMA1001145	F-MAMMA1001145	0.05
	MAMMA1001161	F-MAMMA1001161	0.11
	MAMMA1001162	F-MAMMA1001162	0.11
	MAMMA1001186	F-MAMMA1001186	0.09
. 10	MAMMA1001191	F-MAMMA1001191	0.08
	MAMMA1001203	F-MAMMA1001203	0.08
	MAMMA1001206	F-MAMMA1001206	0.09
	MAMMA1001220	F-MAMMA1001220	0. 1
15	MAMMA1001243	F-MAMMA1001243	0.14
15	MAMMA1001249	F-MAMMA1001249	0.13
	MAMMA1001256	F-MAMMA1001256,	
	MAMMA1001268	F-MAMMA1001268	0, 26
	MAMMA1001274	F-MAMMA1001274	0.13
20	MAMMA1001280	F-MAMMA1001280	0. 1
	MAMMA1001298	F-MAMMA1001298	0.07
	MAMMA1001322	F-MAMMA1001322	0.18
	MAMMA1001324	F-MAMMA1001324	0. 25
25	MAMMA1001330	F-MAMMA1001330	0. 13
	MAMMA1001341	F-MAMMA1001341	0.16
	MAMMA1001383	F-MAMMA1001383	0.05
	MAMMA1001397	F-MAMMA1001397	0. 22
30	MAMMA1001408	F-MAMMA1001408	0.06
	MAMMA1001420	F-MAMMA1001420	0.1
	MAMMA1001435	F-MAMMA1001435	0.07
	MAMMA1001442	F-MAMMA1001442	0.13
35	MAMMA1001446	F-MAMMA1001446	0.08
	MAMMA1001452	F-MAMMA1001452	0.1
	MAMMA1001501	F-MAMMA1001501	0.12
	MAMMA1001502	F-MAMMA1001502	0.09
40	MAMMA1001547	F-MAMMA1001547	0. 22
	MAMMA1001575	F-MAMMA1001575	0.08
	MAMMA1001590	F-MAMMA1001590	0.14
	MAMMA1001606	F-MAMMA1001606	0.13
45	MAMMA1001663	F-MAMMA1001663	0.11
45	MAMMA1001670	F-MAMMA1001670	0. 2
	MAMMA1001671	F-MAMMA1001671	0.13
	MAMMA1001679	F-MAMMA1001679	0.13
	MAMMA1001683	F-MAMMA1001683	0.16
50	MAMMA1001686	F-MAMMA1001686	0.09
	MAMMA1001711	F-MAMMA1001711	0.11
	MAMMA1001715	F-MAMMA1001715	0.11
	MAMMA1001740	F-MAMMA1001740	0.1
55	MAMMA1001745	F-MAMMA1001745	0.09

	MAMMA1001760	F-MAMMA1001760	0.16
	MAMMA1001769	F-MAMMA1001769	0.09
	MAMMA1001788	F-MAMMA1001788	0.1
5	MAMMA1001806	F-MAMMA1001806	0.06
	MAMMA1001815	F-MAMMA1001815	0.1
	MAMMA1001818	F-MAMMA1001818	0.1
	MAMMA1001820	F-MAMMA1001820	0. 27
10	MAMMA1001836	F-MAMMA1001836	0.05
	MAMMA1001854	F-MAMMA1001854	0. 17
	MAMMA1001878	F-MAMMA1001878	0. 13
	MAMMA1001880	F-MAMMA1001880	0. 15
15	MAMMA1001890	F-MAMMA1001890	0.05
	MAMMA1001907	F-MAMMA1001907	0.1
	MAMMA1001908	F-MAMMA1001908,	0.3
	MAMMA1001931	F-MAMMA1001931	0.14
20	MAMMA1001963	F-MAMMA1001963	0.08
	MAMMA1001992	F-MAMMA1001992	0.12
	MAMMA1002032	F-MAMMA1002032	0.08
	MAMMA1002056	F-MAMMA1002056	0.09
25	MAMMA1002058	F-MAMMA1002058	0.16
25	MAMMA1002078	F-MAMMA1002078	0. 28
	MAMMA1002082	F-MAMMA1002082	0.08
	MAMMA1002084	F-MAMMA1002084	0. 24
	MAMMA1002093	F-MAMMA1002093	0.12
30	MAMMA1002108	F-MAMMA1002108	0.05
	MAMMA1002118	F-MAMMA1002118	0.1
	MAMMA1002125	F-MAMMA1002125	0.09
	MAMMA1002132	F-MAMMA1002132	0.11
35	MAMMA1002140	F-MAMMA1002140	0.11
	MAMMA1002145	F-MAMMA1002145	0.11
	MAMMA1002155	F-MAMMA1002155	0.1
	MAMMA1002158	F-MAMMA1002158	0.11
40	MAMMA1002215	F-MAMMA1002215	0.05
	MAMMA1002230	F-MAMMA1002230	0. 07
	MAMMA1002250	F-MAMMA1002250	0. 2
	MAMMA1002267	F-MAMMA1002267	0.08
45	MAMMA1002282	F-MAMMA1002282	0. 05
	MAMMA1002293	F-MAMMA1002293	0.09
	MAMMA1002298	F-MAMMA1002298	0.09
	MAMMA1002299	F-MAMMA1002299	0.16
50	MAMMA1002310	F-MAMMA1002310	0. 16
	MAMMA1002311	F-MAMMA1002311	0. 13
	MAMMA1002322	F-MAMMA1002322	0. 07
	MAMMA1002332	F-MAMMA1002332	0.08
	MAMMA1002339	F-MAMMA1002339	0.09
55	MAMMA1002347	F-MAMMA1002347	0. 06

	MAMMA1002352	F-MAMMA1002352	0.08
	MAMMA1002359	F-MAMMA1002359	0.13
	MAMMA1002360	F-MAMMA1002360	0.17
5	MAMMA1002361	F-MAMMA1002361	0.13
	MAMMA1002392	**	0.16
	MAMMA1002411		0. 28
	MAMMA1002417		0. 19
10	MAMMA1002428	F-MAMMA1002428	0. 07
	MAMMA1002446	F-MAMMA1002446	
	MAMMA1002475	F-MAMMA1002475	
	MAMMA1002480	F-MAMMA1002480	
15	MAMMA1002494	F-MAMMA1002494	
·-	MAMMA1002494	F-MAMMA1002498	0. 17
	MAMMA1002545	F-MAMMA1002545,	
	MAMMA1002556	F-MAMMA1002556	
	MAMMA1002566	F-MAMMA1002566	
20	MAMMA1002571	F-MAMMA1002571	0. 13
	MAMMA1002573	F-MAMMA1002573	0. 29
	MAMMA1002597	F-MAMMA1002597	0. 23
	MAMMA1002603	F-MAMMA1002603	0. 1
25	MAMMA1002612	F-MAMMA1002612	0. 14
	MAMMA1002618	F-MAMMA1002618	0. 11
	MAMMA1002622	F-MAMMA1002622	
	MAMMA1002623		0. 24
30	MAMMA1002625		0. 11
	MAMMA1002629		0. 05
	MAMMA1002636	F-MAMMA1002636	0. 12
	MAMMA1002646	F-MAMMA1002646	0. 05
35	MAMMA1002662		0. 09
	MAMMA1002698		0. 08
	MAMMA1002701		0. 11
	MAMMA1002708		0. 09
40	MAMMA1002721	F-MAMMA1002721	0.06
40	MAMMA1002727	F-MAMMA1002727	0. 15
	MAMMA1002728	F-MAMMA1002728	0. 12
	MAMMA1002744		0. 08
	MAMMA1002746	F-MAMMA1002746	0. 09
45	MAMMA1002748	F-MAMMA1002748	0. 1
	MAMMA1002754	F-MAMMA1002754	0. 07
	MAMMA1002758	F-MAMMA1002758	0. 29
	MAMMA1002764	F-MAMMA1002764	0. 2
50	MAMMA1002765	F-MAMMA1002765	0. 29
	MAMMA1002780	F-MAMMA1002780	0. 12
	MAMMA1002820	F-MAMMA1002820	0. 12
	MAMMA1002830	F-MAMMA1002830	0. 16
55	MAMMA1002833	F-MAMMA1002833	0. 14
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	MAMMA1002835	F-MAMMA1002835	0. 13
	MAMMA1002838	F-MAMMA1002838	0. 09
	MAMMA1002844	F-MAMMA1002844	0. 27
5	MAMMA1002858	F-MAMMA1002858	
	MAMMA1002871	F-MAMMA1002871	0.11
	MAMMA1002870	F-MAMMA1002870	0.11
	MAMMA1002887	F-MAMMA1002887	0. 11
10	MAMMA1002892	F-MAMMA1002892	0.09
	MAMMA1002998	F-MAMMA1002992	0.17
	MAMMA1002908	F-MAMMA1002909	0. 11
	MAMMA1002909	F-MAMMA1002909	0. 22
15	MAMMA1002930	F-MAMMA1002930	0. 2
,,	MAMMA1002941	F-MAMMA1002941	0. 23
	MAMMA1002970	F-MAMMA1002970	
	MAMMA1002972	F-MAMMA1002972,	0. 08 0. 1
	MAMMA1002973	F-MAMMA1002973	0.1
20	MAMMA1003003	F-MAMMA1003003	0.1
	MAMMA1003003	F-MAMMA1003003	0. 2
	MAMMA1003004	F-MAMMA1003004	0.14
	MAMMA1003007	F-MAMMA1003007	0.09
25	MAMMA1003019	F-MAMMA1003019	0.12
	MAMMA1003031	F-MAMMA1003031	0.18
	MAMMA1003039	F-MAMMA1003039	0. 18
	MAMMA1003040	F-MAMMA1003040	0.03
30	MAMMA1003055	F-MAMMA1003055	0.08
	MAMMA1003089	F-MAMMA1003089	0.14
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35	NT2RM4000046	F-NT2RM4000046	0.1
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	NT2RM4000086	F-NT2RM4000086	0. 28
	NT2RM4000200	F-NT2RM4000200	0.16
40	NT2RM4000244	F-NT2RM4000244	0.07
	NT2RM4000265	F-NT2RM4000265	0.18
	NT2RM4000327	F-NT2RM4000327	0.06
	NT2RM4000366	F-NT2RM4000366	0. 28
	NT2RM4000368	F-NT2RM4000368	0.11
45	NT2RM4000414	F-NT2RM4000414	0.19
	NT2RM4000425	F-NT2RM4000425	0.22
	NT2RM4000532	F-NT2RM4000532	0.14
	NT2RM4000779	F-NT2RM4000779	0.13
50	NT2RM4000855	F-NT2RM4000855	0.13
	NT2RM4000979	F-NT2RM4000979	0.22
	NT2RM4001016	F-NT2RM4001016	0.2
	NT2RM4001160	F-NT2RM4001160	0.1
55	NT2RM4001313	F-NT2RM4001313	0. 26

	NT2RM4001414	F-NT2RM4001414	0. 23
	NT2RM4001437	F-NT2RM4001437	0.12
5	NT2RM4001519	F-NT2RM4001519	0. 07
3	NT2RM4001557	F-NT2RM4001557	0.3
	NT2RM4001605	F-NT2RM4001605	0. 22
	NT2RM4001754	F-NT2RM4001754	0. 26
	NT2RM4001776	F-NT2RM4001776	0.18
10	NT2RM4001810	F-NT2RM4001810	0. 22
	NT2RM4001856	F-NT2RM4001856	0.11
	NT2RM4001858	F-NT2RM4001858	0. 14
	NT2RM4001930	F-NT2RM4001930	0. 3
15	NT2RM4001953	F-NT2RM4001953	0. 09
	NT2RM4001984	F-NT2RM4001984	0. 09
	NT2RM4002067	F-NT2RM4002067,	
	NT2RM4002278	F-NT2RM4002278	
20	NT2RM4002278	F-NT2RM4002278	0. 07
20	NT2RM4002287		
		F-NT2RM4002287	0. 27
	NT2RM4002383 NT2RM4002390	F-NT2RM4002383	0.08
		F-NT2RM4002390	0. 18
25	NT2RM4002438	F-NT2RM4002438	0. 23
	NT2RM4002479	F-NT2RM4002479	0. 27
	NT2RM4002499	F-NT2RM4002499	0.08
	NT2RM4002504	F-NT2RM4002504	0.08
30	NT2RM4002532	F-NT2RM4002532	0. 15
	NT2RM4002567	F-NT2RM4002567	0. 28
	NT2RP2000027	F-NT2RP2000027	0.07
	NT2RP2000076	F-NT2RP2000076	0.16
	NT2RP2000077	F-NT2RP2000077	0. 2
35	NT2RP2000098	F-NT2RP2000098	0. 23
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	NT2RP2000183	F-NT2RP2000183	0. 12
	NT2RP2000257	F-NT2RP2000257	0. 27
40	NT2RP2000258	F-NT2RP2000258	0. 19
	NT2RP2000270	F-NT2RP2000270	0.06
	NT2RP2000289	F-NT2RP2000289	0.06
	NT2RP2000327	F-NT2RP2000327	0.08
45	NT2RP2000337	F-NT2RP2000337	0. 14
	NT2RP2000420	F-NT2RP2000420	0. 2
	NT2RP2000459	F-NT2RP2000459	0.12
	NT2RP2000498	F-NT2RP2000498	0.05
•	NT2RP2000523	F-NT2RP2000523	0.14
50	NT2RP2000603	F-NT2RP2000603	0. 22
	NT2RP2000644	F-NT2RP2000644	0.05
	NT2RP2000678	F-NT2RP2000678	0. 11
	NT2RP2000715	F-NT2RP2000715	0. 21
55	NT2RP2000731	F-NT2RP2000731	0. 09
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	NT2RP2000842	F-NT2RP2000758	0.2
		F-NT2RP2000842	0.14
5	NT2RP2000970	F-NT2RP2000970	0. 12
	NT2RP2000987	F-NT2RP2000987	~
	NT2RP2001149	F-NT2RP2001149	0.08
	NT2RP2001196	F-NT2RP2001196	0. 27
	NT2RP2001226	F-NT2RP2001226	0. 18
10	NT2RP2001277	F-NT2RP2001277	0.09
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	NT2RP2001295	F-NT2RP2001295	0.13
	NT2RP2001312	F-NT2RP2001312	0. 15
15	NT2RP2001347	F-NT2RP2001347	0.11
	NT2RP2001423	F-NT2RP2001423	0.06
	NT2RP2001445	F-NT2RP2001445,	0.11
	NT2RP2001449	F-NT2RP2001449	0.15
20	NT2RP2001467	F-NT2RP2001467	0. 21
	NT2RP2001506	F-NT2RP2001506	0.06
	NT2RP2001526	F-NT2RP2001526	0. 1
	NT2RP2001569	F-NT2RP2001569	0.09
25	NT2RP2001663	F-NT2RP2001663	0.08
25	NT2RP2001677	F-NT2RP2001677	0.14
	NT2RP2001678	F-NT2RP2001678	0.17
	NT2RP2001699	F-NT2RP2001699	0. 05
	NT2RP2001720	F-NT2RP2001720	0.08
30	NT2RP2001740	F-NT2RP2001740	0. 23
	NT2RP2001762	F-NT2RP2001762	0.11
	NT2RP2001813	F-NT2RP2001813	0. 19
	NT2RP2001869	F-NT2RP2001869	0. 22
35	NT2RP2001926	F-NT2RP2001926	0. 07
	NT2RP2001936	F-NT2RP2001936	0. 08
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40	NT2RP2002041	F-NT2RP2002041	0. 1
	NT2RP2002047	F-NT2RP2002047	0.12
	NT2RP2002066	F-NT2RP2002066	0. 25
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	NT2RP2002105	F-NT2RP2002105	0.13
45	NT2RP2002124	F-NT2RP2002124	0.13
	NT2RP2002137	F-NT2RP2002137	0.07
	NT2RP2002172	F-NT2RP2002172	0. 21
	NT2RP2002192	F-NT2RP2002192	0. 05
50	NT2RP2002219	F-NT2RP2002219	0. 13
	NT2RP2002256	F-NT2RP2002256	0.17
	NT2RP2002259	F-NT2RP2002259	0. 28
	NT2RP2002316	F-NT2RP2002316	0. 15
55	NT2RP2002394	F-NT2RP2002394	0.16
			2

	NT2RP2002439	F-NT2RP2002439	0. 07
	NT2RP2002457	F-NT2RP2002457	0, 11
	NT2RP2002475	F-NT2RP2002475	0.14
5	NT2RP2002504	F-NT2RP2002504	
	NT2RP2002546	F-NT2RP2002546	0.17
	NT2RP2002591	F-NT2RP2002591	0. 15
	NT2RP2002606	F-NT2RP2002606	0.11
10	NT2RP2002643	F-NT2RP2002643	0. 26
	NT2RP2002727	F-NT2RP2002727	0. 26
	NT2RP2002736	F-NT2RP2002736	0.12
	NT2RP2002740	F-NT2RP2002740	0.16
15	NT2RP2002741	F-NT2RP2002741	0.11
	NT2RP2002750	F-NT2RP2002750	0.14
	NT2RP2002752	F-NT2RP2002752	
	NT2RP2002753	F-NT2RP2002753	0.09
20	NT2RP2002778	F-NT2RP2002778	0.13
20	NT2RP2002839	F-NT2RP2002839	
	NT2RP2002857	F-NT2RP2002857	0. 21
	NT2RP2002987	F-NT2RP2002987	0.08
	NT2RP2003000	F-NT2RP2003000	0. 24
25	NT2RP2003073	F-NT2RP2003073	0.08
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	NT2RP2003161	F-NT2RP2003161	0.05
	NT2RP2003164	F-NT2RP2003164	0.09
30	NT2RP2003206	F-NT2RP2003206	0.22
	NT2RP2003230	F-NT2RP2003230	0.08
	NT2RP2003237	F-NT2RP2003237	0. 24
	NT2RP2003339	F-NT2RP2003339	0. 1
35	NT2RP2003394	F-NT2RP2003394	0.07
	NT2RP2003456	F-NT2RP2003456	0.08
	NT2RP2003499	F-NT2RP2003499	0.08
	NT2RP2003517	F-NT2RP2003517	0. 29
40	NT2RP2003522	F-NT2RP2003522	0. 22
	NT2RP2003559	F-NT2RP2003559	0. 12
	NT2RP2003668	F-NT2RP2003668	0.08
	NT2RP2003704	F-NT2RP2003704	0.09
45	NT2RP2003706	F-NT2RP2003706	0. 28
45	NT2RP2003727	F-NT2RP2003727	0.17
	NT2RP2003770	F-NT2RP2003770	0.09
	NT2RP2003859	F-NT2RP2003859	0.18
	NT2RP2003871	F-NT2RP2003871	0. 23
50	NT2RP2003885	F-NT2RP2003885	0. 24
	NT2RP2003912	F-NT2RP2003912	0. 26
	NT2RP2003968	F-NT2RP2003968	0. 2
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55	NT2RP2004014	F-NT2RP2004014	0. 07

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	NT2RP2004170		0.09
	NT2RP2004172		0.15
5	NT2RP2004207		
	NT2RP2004232		0.17
	NT2RP2004270	F-NT2RP2004270	0. 29
	NT2RP2004300		0.11
10	NT2RP2004321	F-NT2RP2004321	0. 13
	NT2RP2004339		0. 14
	NT2RP2004347		0.14
	NT2RP2004396	F-NT2RP2004396	0. 29
15	NT2RP2004400		0.06
	NT2RP2004412		0. 24
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	NT2RP2004490		0.13
20	NT2RP2004512		0. 1
	NT2RP2004523		0. 15
	NT2RP2004580		0.12
	NT2RP2004587		0.08
25	NT2RP2004675	F-NT2RP2004675	0.14
25	NT2RP2004681	F-NT2RP2004681	0. 17
	NT2RP2004709		0.08
	NT2RP2004736	F-NT2RP2004736	0.13
	NT2RP2004767	F-NT2RP2004767	0. 21
30	NT2RP2004775	F-NT2RP2004775	0.17
	NT2RP2004961	F-NT2RP2004961	0.21
	NT2RP2004962	F-NT2RP2004962	0. 24
	NT2RP2004967	F-NT2RP2004967	0.05
35	NT2RP2004982	F-NT2RP2004982	0.11
	NT2RP2005020	F-NT2RP2005020	0.16
	NT2RP2005031	F-NT2RP2005031	0.07
	NT2RP2005108	F-NT2RP2005108	0.11
40	NT2RP2005254	F-NT2RP2005254	0.16
	NT2RP2005289	F-NT2RP2005289	0. 22
	NT2RP2005293	F-NT2RP2005293	0.08
	NT2RP2005325	F-NT2RP2005325	
45	NT2RP2005336	F-NT2RP2005336	0. 27
	NT2RP2005344	F-NT2RP2005344	0. 24
	NT2RP2005354	F-NT2RP2005354	0.08
	NT2RP2005360	F-NT2RP2005360	0. 12
50	NT2RP2005407	F-NT2RP2005407	0. 28
~ <b>~</b>	NT2RP2005476	F-NT2RP2005476	0.14
	NT2RP2005491	F-NT2RP2005491	0.18
	NT2RP2005501	F-NT2RP2005501	0. 1
	NT2RP2005531	F-NT2RP2005531	0. 29
55	NT2RP2005581	F-NT2RP2005581	0. 07

	NT2RP2005645	F-NT2RP2005645	0. 13
	NT2RP2005651	F-NT2RP2005651	0.06
	NT2RP2005694	F-NT2RP2005694	0.3
5	NT2RP2005701	F-NT2RP2005701_	0.11
	NT2RP2005719	F-NT2RP2005719	0.06
	NT2RP2005726	F-NT2RP2005726	0. 29
	NT2RP2005741	F-NT2RP2005741	0. 22
10	NT2RP2005753	F-NT2RP2005753	0.16
	NT2RP2005815	F-NT2RP2005815	0. 18
	NT2RP2005841	F-NT2RP2005841	0. 05
	NT2RP2005857	F-NT2RP2005857	0. 28
	NT2RP2005908	F-NT2RP2005908	0. 05
15	NT2RP2005942	F-NT2RP2005942	0. 26
	NT2RP2005980	F-NT2RP2005980	
	NT2RP2006098	F-NT2RP2006098	0. 06
	NT2RP2006103	F-NT2RP2006103	0. 07
20	NT2RP2006166	F-NT2RP2006166	0. 07
	NT2RP2006184	F-NT2RP2006184	0. 23
	NT2RP2006258	F-NT2RP2006258	0. 23
	NT2RP2006261	F-NT2RP2006261	0. 12
25	NT2RP2006320	F-NT2RP2006320	0. 16
	NT2RP2006321	F-NT2RP2006321	0. 15
	NT2RP2006323	F~NT2RP2006323	0. 15
	NT2RP2006323	F-NT2RP2006323	0, 07
30	NT2RP2006393	F-NT2RP2006454	0. 07
30	NT2RP2006467	F-NT2RP2006467	0. 11
	NT2RP2006467	F-NT2RP2006467	0. 24
	NT2RP2006554	F-NT2RP2006554	0. 14
	NT2RP2006598	F-NT2RP2006598	0. 16
35	NT2RP3000055	F-NT2RP3000055	0. 08
	NT2RP3000109	F-NT2RP3000109	0. 14
	NT2RP3000109	F-NT2RP3000186	0. 16
			0. 00
40	NT2RP3000233 NT2RP3000235	F-NT2RP3000233 F-NT2RP3000235	0. 29
		,,	
	NT2RP3000247	F-NT2RP3000247	0.19
	NT2RP3000341	F-NT2RP3000341	0. 16
45	NT2RP3000418	F-NT2RP3000418	0. 15
	NT2RP3000433	F-NT2RP3000433	0.11
	NT2RP3000441	F-NT2RP3000441	0. 2
	NT2RP3000449	F-NT2RP3000449	0.16
50	NT2RP3000451	F-NT2RP3000451	0.14
50	NT2RP3000512	F-NT2RP3000512	0.08
	NT2RP3000542	F-NT2RP3000542	0.1
	NT2RP3000561	F-NT2RP3000561	0. 1
	NT2RP3000578	F-NT2RP3000578	0.3
55	NT2RP3000582	F-NT2RP3000582	0.08

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	NT2RP3000584 F-NT2RP3000584	0. 07
	NT2RP3000622 F-NT2RP3000622	0. 17
5	NT2RP3000628 F-NT2RP3000628	0. 08
_	NT2RP3000685 F-NT2RP3000685_	_0, 17
	NT2RP3000736 F-NT2RP3000736	0. 26
	NT2RP3000742 F-NT2RP3000742	0. 22
	NT2RP3000815 F-NT2RP3000815	0.06
10	NT2RP3000865 F-NT2RP3000865	0. 13
	NT2RP3000875 F-NT2RP3000875	0. 16
	NT2RP3000904 F-NT2RP3000904	0. 13
	NT2RP3001007 F-NT2RP3001007	0. 1
15	NT2RP3001115 F-NT2RP3001115	0.1
	NT2RP3001232 F-NT2RP3001232	0.09
	NT2RP3001245 F-NT2RP3001245	0.09
	NT2RP3001281 F-NT2RP3001281	0. 29
20	NT2RP3001318 F-NT2RP3001318	0.12
	NT2RP3001339 F-NT2RP3001339	0. 27
	NT2RP3001340 F-NT2RP3001340	0.11
	NT2RP3001374 F-NT2RP3001374	0. 21
25	NT2RP3001383 F-NT2RP3001383	0.09
25	NT2RP3001432 F-NT2RP3001432	0. 29
	NT2RP3001459 F-NT2RP3001459	0. 16
	NT2RP3001527 F-NT2RP3001527	0.3
	NT2RP3001580 F-NT2RP3001580	0. 23
30	NT2RP3001589 F-NT2RP3001589	0.12
	NT2RP3001607 F-NT2RP3001607	0.12
	NT2RP3001634 F-NT2RP3001634	0. 15
	NT2RP3001752 F-NT2RP3001752	0.09
35	NT2RP3001782 F-NT2RP3001782	0. 21
	NT2RP3001898 F-NT2RP3001898	0.07
	NT2RP3001926 F-NT2RP3001926	0. 05
	NT2RP3001989 F-NT2RP3001989	0.14
40	NT2RP3002002 F-NT2RP3002002	0. 18
	NT2RP3002004 F-NT2RP3002004	0.11
	NT2RP3002056 F-NT2RP3002056	0.16
	NT2RP3002057 F-NT2RP3002057	0.08
45	NT2RP3002166 F-NT2RP3002166	0.15
45	NT2RP3002173 F-NT2RP3002173	0. 28
	NT2RP3002352 F-NT2RP3002352	0.19
	NT2RP3002687 F-NT2RP3002687	0.12
	NT2RP3002713 F-NT2RP3002713	0.1
50	NT2RP3002770 F-NT2RP3002770	0. 23
	NT2RP3002799 F-NT2RP3002799	0.17
	NT2RP3002810 F-NT2RP3002810	0. 2
	NT2RP3002955 F-NT2RP3002955	0. 17
55	NT2RP3002978 F-NT2RP3002978	0.06

	NT2RP3003032 F-NT2RP3003032	0. 1
	NT2RP3003059 F-NT2RP3003059	0.11
	NT2RP3003121 F-NT2RP3003121	0. 23
5	NT2RP3003133 F-NT2RP3003133	
	NT2RP3003138 F-NT2RP3003138	0. 29
	NT2RP3003264 F-NT2RP3003264	0. 15
	NT2RP3003346 F-NT2RP3003346	0. 16
10	NT2RP3003384 F-NT2RP3003384	0. 12
	NT2RP3003403 F-NT2RP3003403	0. 26
	NT2RP3003433 F-NT2RP3003433	0. 22
	NT2RP3003464 F-NT2RP3003464	0. 28
15	NT2RP3003500 F-NT2RP3003500	0. 2
,,	NT2RP3003572 F-NT2RP3003572	0. 29
	NT2RP3003576 F-NT2RP3003576,	
	NT2RP3003625 F-NT2RP3003625	0. 29
	NT2RP3003665 F-NT2RP3003665	0. 15
20	NT2RP3003746 F-NT2RP3003746	0.07
	NT2RP3003799 F-NT2RP3003799	0. 13
	NT2RP3003800 F-NT2RP3003800	0. 28
	NT2RP3003819 F-NT2RP3003819	0. 19
25	NT2RP3003828 F-NT2RP3003828	0. 12
	NT2RP3003842 F-NT2RP3003842	0.11
	NT2RP3003989 F-NT2RP3003989	0.1
	NT2RP3004016 F-NT2RP3004016	0.09
30	NT2RP3004070 F-NT2RP3004070	0.11
	NT2RP3004095 F-NT2RP3004095	0.1
	NT2RP3004110 F-NT2RP3004110	0.07
	NT2RP3004145 F-NT2RP3004145	0. 15
35	NT2RP3004148 F-NT2RP3004148	0.14
	NT2RP3004215 F-NT2RP3004215	0.16
	NT2RP3004334 F-NT2RP3004334	0.1
	NT2RP3004349 F-NT2RP3004349	0.09
40	NT2RP3004399 F-NT2RP3004399	0.16
	NT2RP3004470 F-NT2RP3004470	0. 23
		0. 26
	NT2RP3004503 F-NT2RP3004503	0.12
45		0.06
45	NT2RP4000023 F-NT2RP4000023	0.13
		0.17
		0. 1
		0. 28
50		0. 15
		0. 19
		0. 12
		0. 17
55	NT2RP4000321 F-NT2RP4000321	0. 18

	NT2RP4000424	F-NT2RP4000424	0. 22
	NT2RP4000515	F-NT2RP4000515	0.13
5	NT2RP4000517	F-NT2RP4000517	0.1
	NT2RP4000519	F-NT2RP4000519	0. 26
	NT2RP4000915	F-NT2RP4000915	0.19
	NT2RP4000996	F-NT2RP4000996	0.05
40	NT2RP4001407	F-NT2RP4001407	0. 2
10	NT2RP4001889	F-NT2RP4001889	0.09
	NT2RP4002075	F-NT2RP4002075	0.19
	NT2RP4002905	F-NT2RP4002905	0.15
	OVARC1000017	F-0VARC1000017	0.12
15	0VARC1000058	F-0VARC1000058	0.2
	0VARC1000068	F-0VARC1000068	0.14
	OVARC1000071	F-0VARC1000071,	0.15
	OVARC1000085	F-0VARC1000085	0.16
20	OVARC1000092	F-0VARC1000092	0. 26
	OVARC1000133	F-0VARC1000133	0.14
	OVARC1000145	F-0VARC1000145	0.15
	OVARC1000191	F-0VARC1000191	0.07
25	OVARC1000198	F-0VARC1000198	0. 23
	0VARC1000240	F-0VARC1000240	0.11
	0VARC1000302	F~0VARC1000302	0.08
	0VARC1000414	F-0VARC1000414	0.12
30	0VARC1000427	F-0VARC1000427	0.1
	0VARC1000431	F-0VARC1000431	0.06
	0VARC1000486	F-0VARC1000486	0.08
	0VARC1000496	F-0VARC1000496	0. 13
35	0VARC1000526	F-0VARC1000526	0.09
35	0VARC1000533	F-0VARC1000533	0.11
	0VARC1000543	F-0VARC1000543	0. 2
	0VARC1000573	F-0VARC1000573	0.09
	0VARC1000578	F-0VARC1000578	0. 13
40	0VARC1000678	F-0VARC1000678	0. 17
	0VARC1000679	F-0VARC1000679	0. 23
	0VARC1000700	F-0VARC1000700	0. 28
	0VARC1000769	F-0VARC1000769	0.05
45	OVARC1000802	F-0VARC1000802	0.06
	0VARC1000891	F-0VARC1000891	0.08
	0VARC1000897	F-0VARC1000897	0.11
	0VARC1000936	F-0VARC1000936	0.14
50	0VARC1000937 0VARC1000948	F-0VARC1000937 F-0VARC1000948	0. 29
	OVARC1000960 OVARC1000971	F-0VARC1000960	0.07
	0VARC1000971	F-0VARC1000971 F-0VARC1001000	0.11
			0.06
	OVARC1001011	F-0VARC1001011	0. 15

	OVARC1001051 F-0VARC1001051	0. 16
	OVARC1001062 F-0VARC1001062	0. 09
-	OVARC1001072 F-0VARC1001072	0. 11
5	OVARC1001085 F-0VARC1001085	0. 12
	OVARC1001117 F-0VARC1001117	0. 21
	OVARC1001118 F-0VARC1001118	0. 08
	OVARC1001129 F-0VARC1001129	0. 28
10	OVARC1001169 F-0VARC1001169	0. 12
	OVARC1001240 F-0VARC1001240	0. 09
	OVARC1001261 F-0VARC1001261	0. 1
	OVARC1001268 F-0VARC1001268	0. 05
15	OVARC1001282 F-0VARC1001282	0. 09
		0. 08
	OVARC1001339 F-0VARC1001339, (	0. 15
	3	0. 16
20		0. 06
	0VARC1001442 F-0VARC1001442	0. 15
	OVARC1001480 F-OVARC1001480 (	0. 23
	OVARC1001542 F-OVARC1001542 (	0. 13
	OVARC1001547 F-OVARC1001547	0. 12
25	OVARC1001600 F-0VARC1001600 (	0. 13
	OVARC1001615 F-OVARC1001615 (	). 11
	OVARC1001668 F-OVARC1001668 (	0. 05
	OVARC1001726 F-OVARC1001726 (	). 18
30	OVARC1001745 F-0VARC1001745 (	). 22
	OVARC1001795 F-OVARC1001795 (	). 28
	OVARC1001805 F-0VARC1001805 (	). 16
	OVARC1001812 F-OVARC1001812 (	0. 05
35	OVARC1001813 F-OVARC1001813 (	). 11
	OVARC1001846 F-0VARC1001846 (	). 13
	OVARC1001883 F-OVARC1001883 (	0. 05
	OVARC1001900 F-OVARC1001900 (	). 28
40	OVARC1001901 F-0VARC1001901 (	0.09
	OVARC1001911 F-0VARC1001911 (	0. 06
	OVARC1001928 F-0VARC1001928 (	). 1
		). 07
45		0.06
45	OVARC1002066 F-0VARC1002066 (	0. 08
	OVARC1002143 F-OVARC1002143 (	). 27
		0. 09
		). 15
50	PLACE1000078 F-PLACE1000078 (	). 12
		. 16
		. 15
		. 05
55	PLACE1000332 F-PLACE1000332 (	). 1

	PLACE1000421	F-PLACE1000421	0.14
	PLACE1000424	F-PLACE1000424	0. 08
	PLACE1000481	F-PLACE1000481	0. 12
5	PLACE1000540	F-PLACE1000540	0. 12
	PLACE1000583	F-PLACE1000583	0.11
	PLACE1000599	F-PLACE1000589	0. 1
	PLACE1000399	F-PLACE1000399	0. 22
10			
	PLACE1000841	F-PLACE1000841	0.09
	PLACE1000931	F-PLACE1000931	0. 22
	PLACE1000979	F-PLACE1000979	0. 16
	PLACE1001007	F-PLACE1001007	0. 19
15	PLACE1001015	F-PLACE1001015	0. 25
	PLACE1001076	F-PLACE1001076	0.08
	PLACE1001088	F-PLACE1001088	
	PLACE1001118	F-PLACE1001118	0. 24
20	PLACE1001136	F-PLACE1001136	0.12
	PLACE1001272	F-PLACE1001272	0.07
	PLACE1001279	F-PLACE1001279	0.08
	PLACE1001323	F-PLACE1001323	0.18
25	PLACE1001377	F-PLACE1001377	0.16
	PLACE1001384	F-PLACE1001384	0.06
	PLACE1001395	F-PLACE1001395	0.08
	PLACE1001414	F-PLACE1001414	0.11
	PLACE1001440	F-PLACE1001440	0. 14
30	PLACE1001468	F-PLACE1001468	0. 05
	PLACE1001502	F-PLACE1001502	0.06
	PLACE1001517	F-PLACE1001517	0.09
	PLACE1001534	F-PLACE1001534	0.08
35	PLACE1001603	F-PLACE1001603	0.08
	PLACE1001611	F-PLACE1001611	0. 25
	PLACE1001640	F-PLACE1001640	0.09
	PLACE1001672	F-PLACE1001672	0.11
40	PLACE1001691	F-PLACE1001691	0.11
	PLACE1001705	F-PLACE1001705	0.1
	PLACE1001716	F-PLACE1001716	0. 15
	PLACE1001720	F-PLACE1001720	0.18
45	PLACE1001745	F-PLACE1001745	0.09
40	PLACE1001746	F-PLACE1001746	0.12
	PLACE1001748	F-PLACE1001748	0.08
	PLACE1001756	F-PLACE1001756	0.17
	PLACE1001799	F-PLACE1001799	0.14
50	PLACE1001821	F-PLACE1001821	0.09
	PLACE1001845	F-PLACE1001845	0.08
	PLACE1001897	F-PLACE1001897	0. 15
	PLACE1002052	F-PLACE1002052	0.13
55	PLACE1002066	F-PLACE1002066	0. 07

	PLACE1002150	E-DI ACE10001E0	0.00
	PLACE1002157	F-PLACE1002150 F-PLACE1002157	0.08
	PLACE1002137 PLACE1002205		0. 15
5	PLACE10022203 PLACE1002227	F-PLACE1002205	0. 13
		F-PLACE1002227	0.12
	PLACE1002256	F-PLACE1002256	0.11
	PLACE1002259	F-PLACE1002259	0.09
10	PLACE1002319	F-PLACE1002319	0. 27
	PLACE1002399	F-PLACE1002399	0.08
	PLACE1002477	F-PLACE1002477	0.17
	PLACE1002493	F-PLACE1002493	0.09
	PLACE1002500	F-PLACE1002500	0. 25
15	PLACE1002514	F-PLACE1002514	0.06
	PLACE1002537	F-PLACE1002537	0. 17
	PLACE1002578	F-PLACE1002578,	0.11
	PLACE1002583	F-PLACE1002583	0. 25
20	PLACE1002591	F~PLACE1002591	0. 28
	PLACE1002598	F-PLACE1002598	0. 29
	PLACE1002604	F-PLACE1002604	0. 15
	PLACE1002768	F-PLACE1002768	0.11
25	PLACE1002772	F-PLACE1002772	0. 15
	PLACE1002782	F-PLACE1002782	0.16
	PLACE1002853	F-PLACE1002853	0.16
	PLACE1002881	F-PLACE1002881	0.09
	PLACE1002962	F-PLACE1002962	0.13
30	PLACE1002968	F-PLACE1002968	0. 23
	PLACE1003153	F-PLACE1003153	0.07
	PLACE1003205	F-PLACE1003205	0. 2
	PLACE1003238	F-PLACE1003238	0. 22
35	PLACE1003258	F-PLACE1003258	0.17
	PLACE1003343	F-PLACE1003343	0.07
	PLACE1003361	F-PLACE1003361	0.19
	PLACE1003373	F-PLACE1003373	0.08
40	PLACE1003375	F-PLACE1003375	0.26
-	PLACE1003401	F-PLACE1003401	0.11
	PLACE1003420	F-PLACE1003420	0. 22
	PLACE1003478	F-PLACE1003478	0.14
	PLACE1003516	F-PLACE1003516	0.17
45	PLACE1003519	F-PLACE1003519	0.14
	PLACE1003521	F-PLACE1003521	0.17
	PLACE1003528	F-PLACE1003528	0.16
	PLACE1003566	F-PLACE1003566	0.16
50	PLACE1003575	F-PLACE1003575	0.06
	PLACE1003583	F-PLACE1003583	0.08
	PLACE1003584	F-PLACE1003584	0.1
	PLACE1003593	F-PLACE1003593	0.15
55	PLACE1003618	F-PLACE1003618	0. 25

	PLACE1003638	F-PLACE1003638	0.06
	PLACE1003760	F-PLACE1003760	0.12
	PLACE1003768	F-PLACE1003768	0. 13
5	PLACE1003833		0.06
	PLACE1003850	F-PLACE1003850	0. 07
	PLACE1003858	F-PLACE1003858	0. 07
	PLACE1003864	F-PLACE1003864	0. 12
10	PLACE1003900	F-PLACE1003900	0. 12
		F-PLACE1003932	0. 11
	PLACE1003932	F-PLACE1003932	
	PLACE1004118		0.3
	PLACE1004242	F-PLACE1004242	0. 05
15	PLACE1004256	F-PLACE1004256	0. 24
	PLACE1004257	F-PLACE1004257	0.06
	PLACE1004274	F-PLACE1004274,	
	PLACE1004284	F-PLACE1004284	0. 13
20	PLACE1004336		0. 27
	PLACE1004384		0. 17
	PLACE1004425		0. 12
	PLACE1004467	F-PLACE1004467	0. 2
25	PLACE1004471	F-PLACE1004471	0.12
25	PLACE1004491	F-PLACE1004491	0.08
	PLACE1004506	F-PLACE1004506	
	PLACE1004518	F-PLACE1004518	0.14
	PLACE1004658	F-PLACE1004658	0. 05
30	PLACE1004681	F-PLACE1004681	0. 13
	PLACE1004693	F-PLACE1004693	0.11
	PLACE1004716	F-PLACE1004716	0.12
	PLACE1004773	F-PLACE1004773	0. 17
35	PLACE1004793	F-PLACE1004793	0, 21
	PLACE1004813	F-PLACE1004813	0.11
	PLACE1004815	F-PLACE1004815	0.14
	PLACE1004827	F-PLACE1004827	0. 15
	PLACE1004836	F-PLACE1004836	0. 25
40	PLACE1004838	F-PLACE1004838	0. 16
	PLACE1004840	F-PLACE1004840	0.06
	PLACE1004900	F-PLACE1004900	0. 08
	PLACE1004913	F-PLACE1004913	0. 12
45	PLACE1004934	F-PLACE1004934	0. 12
	PLACE1004934	F-PLACE1004972	0. 17
			0. 07
	PLACE1004979	F-PLACE1004979	
50	PLACE1004985	F-PLACE1004985	0.05
50	PLACE1005052	F-PLACE1005052	0. 17
	PLACE1005077	F-PLACE1005077	0.1
	PLACE1005085	F-PLACE1005085	0. 13
	PLACE1005086	F-PLACE1005086	0. 13
55	PLACE1005108	F-PLACE1005108	0. 28

	PLACE1005111 F-PLACE1005111	0.15
	PLACE1005128 F-PLACE1005128	0. 12
	PLACE1005146 F-PLACE1005146	0. 07
5	PLACE1005162 F-PLACE1005162	0. 13
	PLACE1005176 F-PLACE1005176	0. 16
	PLACE1005409 F-PLACE1005409	0. 09
	PLACE1005453 F-PLACE1005453	0. 1
10	PLACE1005467 F-PLACE1005467	0. 28
	PLACE1005471 F-PLACE1005471	0.06
	PLACE1005477 F-PLACE1005477	0. 12
	PLACE1005502 F-PLACE1005502	0. 12
	PLACE1005502 F-PLACE1005502 PLACE1005526 F-PLACE1005526	0. 1
15		0.07
	PLACE1005528 F-PLACE1005528	0.07
	PLACE1005574 F-PLACE1005574,	
	PLACE1005584 F-PLACE1005584	0. 24
20	PLACE1005611 F-PLACE1005611	0. 18
	PLACE1005639 F-PLACE1005639	0.09
	PLACE1005666 F-PLACE1005666	0.07
	PLACE1005730 F-PLACE1005730	0. 2
25	PLACE1005802 F-PLACE1005802	0. 1
	PLACE1005845 F-PLACE1005845	0.09
	PLACE1005850 F-PLACE1005850	0. 22
	PLACE1005884 F-PLACE1005884	0.08
	PLACE1005898 F-PLACE1005898	0. 12
30	PLACE1005932 F-PLACE1005932	0. 13
	PLACE1005968 F-PLACE1005968	0. 07
	PLACE1006002 F-PLACE1006002	0.08
	PLACE1006003 F-PLACE1006003	0. 16
35	PLACE1006017 F-PLACE1006017	0. 13
	PLACE1006076 F-PLACE1006076	0.11
	PLACE1006129 F-PLACE1006129	0. 25
	PLACE1006143 F-PLACE1006143	0.08
40	PLACE1006164 F-PLACE1006164	0. 05
	PLACE1006187 F-PLACE1006187	0. 2
	PLACE1006205 F-PLACE1006205	0.08
	PLACÉ1006223 F-PLACE1006223	0. 05
	PLACE1006262 F-PLACE1006262	0.08
45	PLACE1006288 F-PLACE1006288	0.06
	PLACE1006318 F-PLACE1006318	0.12
	PLACE1006360 F-PLACE1006360	0. 07
	PLACE1006368 F-PLACE1006368	0. 27
50	PLACE1006371 F-PLACE1006371	0. 14
	PLACE1006382 F-PLACE1006382	0.09
	PLACE1006470 F-PLACE1006470	0.14
	PLACE1006506 F-PLACE1006506	0. 24
55	PLACE1006521 F~PLACE1006521	0.08
J = -		

	PLACE1006534	F-PLACE1006534	0.06
	PLACE1006540	F-PLACE1006540	0.05
	PLACE1006617	F-PLACE1006617	0.05
5	PLACE1006629	F-PLACE1006629	0.2
	PLACE1006640	F-PLACE1006640	0.05
	PLACE1006754	F-PLACE1006754	0.08
	PLACE1006760	F-PLACE1006760	0.05
10	PLACE1006779	F-PLACE1006779	0.17
	PLACE1006792	F-PLACE1006792	0.15
	PLACE1006795	F-PLACE1006795	0. 1
	PLACE1006800	F-PLACE1006800	0. 11
45	PLACE1006805	F-PLACE1006805	0. 12
15	PLACE1006815	F-PLACE1006815	0.14
	PLACE1006860	F-PLACE1006860,	
	PLACE1006867	F-PLACE1006867	0.11
	PLACE1006878	F-PLACE1006878	0. 23
20	PLACE1006904	F-PLACE1006904	0. 13
	PLACE1007045	F-PLACE1007045	0. 08
	PLACE1007097	F-PLACE1007097	0. 13
	PLACE1007111	F-PLACE1007111	0.08
25	PLACE1007132	F-PLACE1007132	0. 07
	PLACE1007140	F-PLACE1007140	0.16
	PLACE1007276	F-PLACE1007276	0.09
	PLACE1007286	F-PLACE1007286	0. 05
30	PLACE1007478	F-PLACE1007478	0. 19
	PLACE1007525	F-PLACE1007525	0. 11
	PLACE1007557	F-PLACE1007557	0. 13
	PLACE1007677	F-PLACE1007677	0.08
35	PLACE1007737	F-PLACE1007737	0. 07
33	PLACE1007743	F-PLACE1007743	0.09
	PLACE1007807	F-PLACE1007807	0.09
	PLACE1007829	F-PLACE1007829	0.09
	PLACE1007852	F-PLACE1007852	0.14
40	PLACE1007866	F-PLACE1007866	0.07
	PLACE1007877	F-PLACE1007877	0. 29
	PLACE1008045	F-PLACE1008045	0.17
	PLACE1008080	F-PLACE1008080	0.05
45	PLACE1008111	F-PLACE1008111	0.11
	PLACE1008181	F-PLACE1008181	0.08
	PLACE1008201	F-PLACE1008201	0.12
	PLACE1008231	F-PLACE1008231	0.08
50	PLACE1008244	F-PLACE1008244	0.06
	PLACE1008330	F-PLACE1008330	0. 13
	PLACE1008331	F-PLACE1008331	0.15
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55	PLACE1008392	F-PLACE1008392	0.09

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	PLACE1008424	F-PLACE1008424	0.06
	PLACE1008426	F-PLACE1008426	0.17
5	PLACE1008437	F-PLACE1008437	0. 2
	PLACE1008455	F-PLACE1008455	0.07
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	PLACE1008621	F-PLACE1008621	0.07
10	PLACE1008625	F-PLACE1008625	0, 12
	PLACE1008630	F-PLACE1008630	0.12
	PLACE1008643	F-PLACE1008643	0.11
	PLACE1008045	F-PLACE1008715	0. 1
	PLACE1008718	F-PLACE1008748	0. 15
15	PLACE1008748	F-PLACE1008757	0. 13
	PLACE1008798	F-PLACE1008798,	0.14
	PLACE1008807	F-PLACE1008807	0. 12
20	PLACE1008851	F-PLACE1008851	
	PLACE1008920	F-PLACE1008920	0.1
	PLACE1008941	F-PLACE1008941	0.14
	PLACE1008947	F-PLACE1008947	0, 05
25	PLACE1009020	F-PLACE1009020	0.14
	PLACE1009039	F-PLACE1009039	0.16
	PLACE1009048	F-PLACE1009048	0.09
	PLACE1009050	F-PLACE1009050	0.09
	PLACE1009150	F-PLACE1009150	0. 12
30	PLACE1009155	F-PLACE1009155	0.08
	PLACE1009172	F-PLACE1009172	0.07
	PLACE1009183	F-PLACE1009183	0.07
	PLACE1009200	F-PLACE1009200	0.08
35	PLACE1009246	F-PLACE1009246	0.08
	PLACE1009308	F-PLACE1009308	0. 23
	PLACE1009398	F-PLACE1009398	0. 26
	PLACE1009410	F-PLACE1009410	0. 15
40	PLACE1009477	F-PLACE1009477	0.18
	PLACE1009493	F-PLACE1009493	0. 23
	PLACE1009539	F-PLACE1009539	0.11
	PLACE1009595	F-PLACE1009595	0.11
	PLACE1009613	F-PLACE1009613	0.08
45	PLACE1009621	F-PLACE1009621	0.19
	PLACE1009637	F-PLACE1009637	0.27
	PLACE1009639	F-PLACE1009639	0.27
	PLACE1009879	F-PLACE1009879	0. 07
50	PLACE1009888	F-PLACE1009888	0.06
	PLACE1009924	F-PLACE1009924	0.09
	PLACE1009925	F-PLACE1009925	0. 21
	PLACE1009935	F-PLACE1009935	0.22
55	PLACE1009947	F-PLACE1009947	0.19

	PLACE1010069	E-DI ACE1010000	0.00
		F-PLACE1010069	0. 22
	PLACE1010083	F-PLACE1010083	0. 26
_	PLACE1010089	F-PLACE1010089	0. 13
5	PLACE1010181	F-PLACE1010181	0.07
	PLACE1010231	F-PLACE1010231	0. 17
	PLACE1010270	F-PLACE1010270	0.08
	PLACE1010341	F-PLACE1010341	0.1
10	PLACE1010562	F-PLACE1010562	0.14
	PLACE1010579	F-PLACE1010579	0.15
	PLACE1010624	F-PLACE1010624	0.13
	PLACE1010628	F-PLACE1010628	0.16
15	PLACE1010631	F-PLACE1010631	0. 23
	PLACE1010662	F-PLACE1010662	0. 24
	PLACE1010702	F-PLACE1010702,	0. 21
	PLACE1010739	F-PLACE1010739	0.13
20	PLACE1010802	F-PLACE1010802	0.17
20	PLACE1010891	F-PLACE1010891	0.09
	PLACE1010896	F-PLACE1010896	0. 22
	PLACE1010916	F-PLACE1010916	0.15
	PLACE1010947	F-PLACE1010947	0.08
25	PLACE1011032	F-PLACE1011032	0.17
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30	PLACE1011143	F-PLACE1011143	0. 23
	PLACE1011165	F-PLACE1011165	0. 05
	PLACE1011185	F-PLACE1011185	0. 17
	PLACE1011203	F-PLACE1011203	0.08
35	PLACE1011296	F-PLACE1011296	0. 13
	PLACE1011340	F-PLACE1011340	0. 05
	PLACE1011375	F-PLACE1011375	0. 07
	PLACE1011419	F-PLACE1011419	0. 15
	PLACE1011452	F-PLACE1011452	0. 13
40	PLACE1011465	F-PLACE1011465	0.14
	PLACE1011492	F-PLACE1011492	0. 26
	PLACE1011503	F-PLACE1011503	0.14
	PLACE1011520	F-PLACE1011520	0. 12
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	PLACE1011576	F-PLACE1011576	0. 29
	PLACE1011641	F-PLACE1011641	0. 07
	PLACE1011643	F-PLACE1011643	0. 27
50	PLACE1011649	F-PLACE1011649	0. 17
	PLACE1011650	F-PLACE1011650	0.13
	PLACE1011719	F-PLACE1011719	0. 15
	PLACE1011729	F-PLACE1011729	0. 23
55	PLACE1011749	F-PLACE1011749	0.12
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	PLACE1011778	F-PLACE1011778	0.07
	PLACE1011874	F-PLACE1011874	0. 07
5	PLACE1011962	F-PLACE1011962	0.06
	PLACE1011964	F-PLACE1011964	0. 13
	PLACE1011995	F-PLACE1011995	0.08
	PLACE2000003	F-PLACE2000003	0.09
10	PLACE2000011	F-PLACE2000011	0. 12
	PLACE2000017	F-PLACE2000017	0. 22
	PLACE2000061	F-PLACE2000061	0. 19
	PLACE2000103	F-PLACE2000103	0. 16
	PLACE2000115	F-PLACE2000115	0. 11
15	PLACE2000113	F-PLACE2000132	0. 24
	PLACE2000136	F-PLACE2000136	
	PLACE2000130	F-PLACE2000140	0. 24
	PLACE2000176	F-PLACE2000176	0. 25
20	PLACE2000170	F-PLACE2000170	0. 16
	PLACE200016	F-PLACE2000187	0. 15
	PLACE2000210	F-PLACE2000210	0. 15
	PLACE2000305	F-PLACE2000305	0.05
25	PLACE2000303	F-PLACE2000317	0. 25
	PLACE2000317	F-PLACE2000335	0. 09
	PLACE2000333	F-PLACE2000342	0. 03
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	PLACE2000455	F-PLACE2000455	0. 15
40	PLACE2000465	F-PLACE2000465	0. 12
	PLACE2000477	F-PLACE2000477	0. 3
	PLACE3000004	F-PLACE3000004	0. 3
	PLACE3000119	F-PLACE3000119	0. 1
45	PLACE3000124	F-PLACE3000124	0. 11
	PLACE3000136	F-PLACE3000136	0. 13
	PLACE3000158	F-PLACE3000158	0.06
	PLACE3000169	F-PLACE3000169	0. 13
50	PLACE3000109	F-PLACE3000199	0. 16
30	PLACE3000207	F-PLACE3000199	0. 11
	PLACE3000207	F-PLACE3000208	0. 07
	PLACE3000218	F-PLACE3000218	0. 15
	PLACE3000220	F-PLACE3000220	0. 05
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PLACE3000271 F-PLACE3000271 0. PLACE3000276 F-PLACE3000276 0. PLACE3000341 F-PLACE3000341 0. PLACE3000362 F-PLACE3000362 0. PLACE3000365 F-PLACE3000365 0. PLACE3000373 F-PLACE3000373 0. PLACE3000399 F-PLACE3000388 0. PLACE3000400 F-PLACE3000399 0. PLACE3000400 F-PLACE3000400 0. PLACE3000401 F-PLACE3000401 0. PLACE3000402 F-PLACE3000402 0. PLACE3000475 F-PLACE3000475, 0. PLACE400093 F-PLACE400093 0. PLACE4000237 F-PLACE4000233 0. PLACE4000247 F-PLACE4000247 0. PLACE4000250 F-PLACE4000250 0. PLACE4000250 F-PLACE4000250 0. PLACE4000320 F-PLACE4000320 0. PLACE4000320 F-PLACE4000320 0. PLACE4000330 F-PLACE4000320 0. PLACE4000344 F-PLACE4000344 0. PLACE4000379 F-PLACE4000379 0. PLACE4000379 F-PLACE4000379 0. PLACE4000379 F-PLACE4000379 0.	08 19 09 22 12 14 08 1 1 17 07 21 19 07 14 17 09 23 21
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PLACE3000341 F-PLACE3000341 0. PLACE3000362 F-PLACE3000362 0. PLACE3000365 F-PLACE3000365 0. PLACE3000373 F-PLACE3000373 0. PLACE3000388 F-PLACE3000388 0. PLACE3000399 F-PLACE3000399 0. PLACE3000400 F-PLACE3000400 0. PLACE3000401 F-PLACE3000401 0. PLACE3000402 F-PLACE3000402 0. PLACE3000405 F-PLACE3000406 0. PLACE3000475 F-PLACE3000406 0. PLACE3000475 F-PLACE400093 0. PLACE4000093 F-PLACE4000100 0. PLACE4000233 F-PLACE4000233 0. PLACE4000247 F-PLACE4000247 0. PLACE4000250 F-PLACE4000250 0. PLACE4000250 F-PLACE4000250 0. PLACE4000320 F-PLACE4000367 0. PLACE4000344 F-PLACE4000344 0. PLACE4000379 F-PLACE4000379 0. PLACE4000379 F-PLACE4000379 0. PLACE4000379 F-PLACE4000379 0.	19 09 22 12 14 08 1 1 17 07 21 19 07 14 17 09 23 21
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PLACE3000373 F-PLACE3000373 0.0 PLACE3000388 F-PLACE3000388 0.0 PLACE3000399 F-PLACE3000399 0.0 PLACE3000400 F-PLACE3000400 0.0 PLACE3000401 F-PLACE3000401 0.0 PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE400093 F-PLACE400093 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000367 F-PLACE4000367 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0	22 12 14 08 1 1 1 7 07 21 19 07 14 17 09 23 21
PLACE3000388 F-PLACE3000388 0.0 PLACE3000399 F-PLACE3000399 0.0 PLACE3000400 F-PLACE3000400 0.0 PLACE3000401 F-PLACE3000401 0.0 PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE400093 F-PLACE400093 0.0 PLACE4000100 F-PLACE4000100 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000250 0.0 PLACE4000330 F-PLACE4000330 0.0 PLACE4000344 F-PLACE4000350 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000401 0.0	12 14 08 1 1 17 07 21 19 07 14 17 09 23 21
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PLACE3000399 F-PLACE3000399 0.0 PLACE3000400 F-PLACE3000400 0.0 PLACE3000401 F-PLACE3000401 0.0 PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE400093 F-PLACE400093 0.0 PLACE4000093 F-PLACE4000093 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000250 0.0 PLACE4000307 F-PLACE4000300 0.0 PLACE4000307 F-PLACE4000307 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000401 0.0	14 08 1 1 17 07 21 19 07 14 17 09 23 21
PLACE3000400 F-PLACE3000400 0.0 PLACE3000401 F-PLACE3000401 0.0 PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE400093 F-PLACE400093 0.0 PLACE4000093 F-PLACE4000100 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000250 0.0 PLACE4000320 F-PLACE4000320 0.0 PLACE4000344 F-PLACE4000344 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000401 F-PLACE4000401 0.0	08 1 17 07 21 19 07 14 17 09 23 21
PLACE3000401 F-PLACE3000401 0.0 PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE400093 F-PLACE400093 0.0 PLACE4000100 F-PLACE4000100 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000250 0.0 PLACE4000320 F-PLACE4000340 0.0 PLACE4000344 F-PLACE4000340 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE40000401 F-PLACE4000401 0.0	1 1 17 07 21 19 07 14 17 09 23 21
PLACE3000402 F-PLACE3000402 0.0 PLACE3000406 F-PLACE3000406 0.0 PLACE3000475 F-PLACE3000475, 0.0 PLACE4000093 F-PLACE4000093 0.0 PLACE4000100 F-PLACE4000100 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000250 0.0 PLACE4000320 F-PLACE4000320 0.0 PLACE4000344 F-PLACE4000344 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000401 F-PLACE4000401 0.0	1 17 07 21 19 07 14 17 09 23 21
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PLACE400093 F-PLACE4000093 0.0 PLACE4000100 F-PLACE4000100 0.0 PLACE4000233 F-PLACE4000233 0.0 PLACE4000247 F-PLACE4000247 0.0 PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000252 0.0 PLACE4000320 F-PLACE4000320 0.0 PLACE4000344 F-PLACE4000344 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000401 F-PLACE4000401 0.0	21 19 07 14 17 09 23 21
PLACE4000100 F-PLACE4000100 0.00 PLACE4000233 F-PLACE4000233 0.00 PLACE4000247 F-PLACE4000247 0.00 PLACE4000250 F-PLACE4000250 0.00 PLACE4000252 F-PLACE4000252 0.00 PLACE4000320 F-PLACE4000320 0.00 PLACE4000344 F-PLACE4000344 0.00 PLACE4000367 F-PLACE4000367 0.00 PLACE4000379 F-PLACE4000379 0.00 PLACE4000401 F-PLACE4000401 0.00	19 07 14 17 09 23 21
PLACE4000233 · F-PLACE4000233 · 0.0 PLACE4000247 F-PLACE4000247 · 0.0 PLACE4000250 F-PLACE4000250 · 0.0 PLACE4000252 F-PLACE4000252 · 0.0 PLACE4000320 F-PLACE4000320 · 0.0 PLACE4000344 F-PLACE4000344 · 0.0 PLACE4000367 F-PLACE4000367 · 0.0 PLACE4000379 F-PLACE4000379 · 0.0 PLACE4000401 F-PLACE4000401 · 0.0	07 14 17 09 23 21
PLACE4000247 F-PLACE4000247 0. PLACE4000250 F-PLACE4000250 0. PLACE4000252 F-PLACE4000252 0. PLACE4000320 F-PLACE4000320 0. PLACE4000344 F-PLACE4000344 0. PLACE4000367 F-PLACE4000367 0. PLACE4000379 F-PLACE4000379 0. PLACE4000401 F-PLACE4000401 0.	14 17 09 23 21
PLACE4000250 F-PLACE4000250 0.0 PLACE4000252 F-PLACE4000252 0.0 PLACE4000320 F-PLACE4000320 0.0 PLACE4000344 F-PLACE4000344 0.0 PLACE4000367 F-PLACE4000367 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000401 F-PLACE4000401 0.0	17 09 23 21
PLACE4000252 F-PLACE4000252 0.0 PLACE4000320 F-PLACE4000320 0.0 PLACE4000344 F-PLACE4000344 0.0 PLACE4000367 F-PLACE4000367 0.0 PLACE4000379 F-PLACE4000379 0.0 PLACE4000401 F-PLACE4000401 0.0	09 23 21
PLACE4000320 F-PLACE4000320 0. PLACE4000344 F-PLACE4000344 0. PLACE4000367 F-PLACE4000367 0. PLACE4000379 F-PLACE4000379 0. PLACE4000401 F-PLACE4000401 0.	23 21
PLACE4000344 F-PLACE4000344 0. PLACE4000367 F-PLACE4000367 0. PLACE4000379 F-PLACE4000379 0. PLACE4000401 F-PLACE4000401 0.	21
PLACE4000367 F-PLACE4000367 0. PLACE4000379 F-PLACE4000379 0. PLACE4000401 F-PLACE4000401 0.	
PLACE4000379 F-PLACE4000379 0. PLACE4000401 F-PLACE4000401 0.	175
PLACE4000401 F-PLACE4000401 0.	14
	13
PLACE4000411 F-PLACE4000411 0.	11
	19
	27
TUVB01000000 E_TUVB01000000 0	09
	09
	13
	21
	13
40	11
THYR01000132 F-THYR01000132 0.	
	13
	12
	12 11
	12 11 16
THYR01000241 F-THYR01000241 0	12 11 16 08
	12 11 16 08 08
50 THYR01000279 F-THYR01000279 0.	12 11 16 08 08 13
THYR01000279 F-THYR01000279 0. THYR01000438 F-THYR01000438 0.	12 11 16 08 08 13
THYR01000279 F-THYR01000279 0. THYR01000438 F-THYR01000438 0. THYR01000452 F-THYR01000452 0.	12 11 16 08 08 13 06 12
THYR01000279 F-THYR01000279 0. THYR01000438 F-THYR01000438 0. THYR01000452 F-THYR01000452 0. THYR01000471 F-THYR01000471 0.	12 11 16 08 08 13

	THYR01000558	F-THYR01000558	0.09
	THYR01000596	F-THYR01000596	0. 14
	THYR01000602	F-THYR01000602	0. 13
5	THYR01000625	F-THYR01000625	0. 2
	THYR01000641	F-THYR01000641	0.06
	THYR01000658	F-THYR01000658	0. 12
	THYR01000699	F-THYR01000699	0.09
10	THYR01000748	F-THYR01000748	0. 3
	THYR01000793	F-THYR01000793	0. 25
	THYR01000796	F-THYR01000796	0. 11
	THYR01000805	F-THYR01000805	0. 28
	THYR01000815	F-THYR01000815	0. 14
15	THYR01000843	F-THYR01000843	0. 13
	THYR01000852	F-THYR01000852,	
	THYR01000865	F-THYR01000865	0. 14
	THYR01000895	F-THYR01000895	0. 12
20 ,	THYR01000974	F-THYR01000974	0. 07
	THYR01000975	F-THYR01000975	0.08
	THYR01001003	F-THYR01001003	0.11
	THYR01001031	F-THYR01001031	0. 1
25	THYR01001062	F-THYR01001062	0. 14
	THYR01001093	F-THYR01001093	0. 13
	THYR01001121	F-THYR01001121	0.14
	THYR01001133	F-THYR01001133	0. 24
30	THYR01001177	F-THYR01001177	
	THYR01001262	F-THYR01001262	0. 2
	THYR01001290	F-THYR01001290	0.06
	THYR01001321	F-THYR01001321	0.09
35	THYR01001322	F-THYR01001322	0.13
	THYR01001365	F-THYR01001365	0.08
	THYR01001401	F-THYR01001401	0.21
	THYR01001411	F-THYR01001411	0.06
40	THYR01001426	F-THYR01001426	0. 1
	THYR01001434	F-THYR01001434	0. 15
	THYR01001480	F-THYR01001480	0. 1
	THYR01001534	F-THYR01001534	0. 26
	THYR01001541	F-THYR01001541	0. 22
45	THYR01001559	F-THYR01001559	0. 15
	THYR01001570	F-THYR01001570	0. 28
	THYR01001573	F-THYR01001573	0.05
	THYR01001595	F-THYR01001595	0.05
50	THYR01001617	F-THYR01001617	0.1
	THYR01001673	F-THYR01001673	0.07
	THYR01001706	F-THYR01001706	0. 15
	THYR01001745	F-THYR01001745	0. 12
55	THYR01001746	F-THYR01001746	0. 1

	THYR01001772 F-THYR01001772 0.08
	THYR01001895 F-THYR01001895 0.19
5	THYR01001907 F-THYR01001907 0.17
3	Y79AA1000033 F-Y79AA1000033 0.18
	Y79AA1000065 F-Y79AA1000065 0.22
	Y79AA1000346 F-Y79AA1000346 0.2
0	Y79AA1000405 F-Y79AA1000405 0.07
	Y79AA1000410 F-Y79AA1000410 0.08
	Y79AA1000538 F-Y79AA1000538 0.06
	Y79AA1000802 F-Y79AA1000802 0.28
5	Y79AA1000805 F-Y79AA1000805 0.2
	Y79AA1000969 F-Y79AA1000969 0.07
	Y79AA1001061 F-Y79AA1001061 0.22
	Y79AA1001167 F-Y79AA1001167, 0.22
	Y79AA1001384 F-Y79AA1001384 0.18
	Y79AA1001594 F-Y79AA1001594 0.22
	Y79AA1001692 F-Y79AA1001692 0.2
	Y79AA1001848 F-Y79AA1001848 0.21
	Y79AA1001875 F-Y79AA1001875 0.09
	Y79AA1002103 F-Y79AA1002103 0.11
	Y79AA1002211 F-Y79AA1002211 0.19
)	Y79AA1002220 F-Y79AA1002220 0.25

# Table 11

574 clones belonging to groups (4), (5), (6), and (7), each having the maximal ATGpr1 score of not more than 0.3 and selected based on the maximal ATGpr2 score, and their maximal ATGpr2 scores

		<del></del>	
10	clone name	name of sequence	maximal ATGpr2 score
15	HEMBA1000180 HEMBA1000251 HEMBA1000280 HEMBA1000282	F-HEMBA1000251 F-HEMBA1000280	0. 37 0. 14 0. 33 0. 36
20	HEMBA1000333 HEMBA1000351 HEMBA1000357 HEMBA1000376	F-HEMBA1000333 F-HEMBA1000351 F-HEMBA1000357 F-HEMBA1000376	0. 59 0. 36 0. 39 0. 53
25	HEMBA1000469 HEMBA1000504 HEMBA1000519 HEMBA1000545	F-HEMBA1000504 F-HEMBA1000519 F-HEMBA1000545	0. 32 0. 34 0. 72 0. 32
30	HEMBA1000557 HEMBA1000575	F-HEMBA1000557 F-HEMBA1000575	0. 35 0. 34

	HEMBA1000604	F-HEMBA1000604	0. 28
	HEMBA1000622	F-HEMBA1000622	0. 32
	HEMBA1000673	F-HEMBA1000673	0. 14
5	HEMBA1000726	F-HEMBA1000726	
	HEMBA1000749	F-HEMBA1000749	0. 35
	HEMBA1000822	F-HEMBA1000822	0. 42
	HEMBA1000876	F-HEMBA1000876	0. 44
10	HEMBA1000934	F-HEMBA1000934	0. 41
	HEMBA1000943	F-HEMBA1000943	0. 34
	HEMBA1000960	F-HEMBA1000960	0.34
	HEMBA1000972	F-HEMBA1000972	0. 44
15	HEMBA1001020	F-HEMBA1001020	0. 31
	HEMBA1001024	F-HEMBA1001024	0. 22
	HEMBA1001026	F-HEMBA1001026,	
	HEMBA1001051	F-HEMBA1001051	0. 36
20	HEMBA1001060	F-HEMBA1001060	0. 34
20	HEMBA1001071	F-HEMBA1001071	0. 57
	HEMBA1001208	F-HEMBA1001208	0. 57
	HEMBA1001226	F-HEMBA1001226	0. 44
25	HEMBA1001319	F-HEMBA1001319	0. 58
25	HEMBA1001383	F-HEMBA1001383	0. 35
	HEMBA1001391	F-HEMBA1001391	0.42
	HEMBA1001411	F-HEMBA1001411	0. 55
	HEMBA1001433	F-HEMBA1001433	0.66
30	HEMBA1001435	F-HEMBA1001435	0.72
	HEMBA1001478	F-HEMBA1001478	0.36
	HEMBA1001522	F-HEMBA1001522	0.31
	HEMBA1001636	F-HEMBA1001636	0.32
35	HEMBA1001651	F-HEMBA1001651	0.54
	HEMBA1001658	F-HEMBA1001658	0. 38
	HEMBA1001709	F-HEMBA1001709	0.55
	HEMBA1001734	F-HEMBA1001734	0. 13
40	HEMBA1001745	F-HEMBA1001745	0.33
	HEMBA1001784	F-HEMBA1001784	0.08
	HEMBA1001791	F-HEMBA1001791	0.48
	HEMBA1001803	F-HEMBA1001803	0. 21
45	HEMBA1001808	F-HEMBA1001808	0. 23
	HEMBA1001835	F-HEMBA1001835	0. 67
	HEMBA1001844	F-HEMBA1001844	0. 46
	HEMBA1001888	F-HEMBA1001888	0. 72
50	HEMBA1001918	F-HEMBA1001918	0. 25
	HEMBA1001940	F-HEMBA1001940	0.40
	HEMBA1001962	F-HEMBA1001962	0.63
	HEMBA1002022	F-HEMBA1002022	0. 32
55	HEMBA1002185	F-HEMBA1002185	0. 33
	HEMBA1002257	F-HEMBA1002257	0. 13

	HEMBA1002348	F-HEMBA1002348	0.39
	HEMBA1002621	F-HEMBA1002621	0.47
	HEMBA1002645	F-HEMBA1002645	0.34
5	HEMBA1002661	F-HEMBA1002661	0. 43
	HEMBA1002728	F-HEMBA1002728	0.45
	HEMBA1002742	F-HEMBA1002742	0.49
	HEMBA1002921	F-HEMBA1002921	0.79
10	HEMBA1002924	F-HEMBA1002924	0. 45
	HEMBA1002934	F-HEMBA1002934	0.34
	HEMBA1002968	F-HEMBA1002968	0. 39
	HEMBA1003037	F-HEMBA1003037	0.43
15	HEMBA1003064	F-HEMBA1003064	0.19
	HEMBA1003071	F-HEMBA1003071	0.48
	HEMBA1003166	F-HEMBA1003166,	0, 32
	HEMBA1003202	F-HEMBA1003202	0. 38
20	HEMBA1003204	F-HEMBA1003204	0.34
	HEMBA1003220	F-HEMBA1003220	0.39
	HEMBA1003229	F-HEMBA1003229	0.43
	HEMBA1003276	F-HEMBA1003276	0.48
25	HEMBA1003296	F-HEMBA1003296	0.36
20	HEMBA1003373	F-HEMBA1003373	0.48
	HEMBA1003531	F-HEMBA1003531	0.46
	HEMBA1003640	F-HEMBA1003640	0.32
	HEMBA1003714	F-HEMBA1003714	0.16
30	HEMBA1003856	F-HEMBA1003856	0.47
	HEMBA1003902	F-HEMBA1003902	0.17
	HEMBA1003908	F-HEMBA1003908	0.44
	HEMBA1003926	F-HEMBA1003926	0.36
35	HEMBA1003942	F-HEMBA1003942	0.47
	HEMBA1003987	F-HEMBA1003987	0.39
	HEMBA1004000	F-HEMBA1004000	0. 87
	HEMBA1004024	F-HEMBA1004024	0.82
40	HEMBA1004164	F-HEMBA1004164	0. 21
	HEMBA1004267	F-HEMBA1004267	0.46
	HEMBA1004323	F-HEMBA1004323	0.37
	HEMBA1004405	F-HEMBA1004405	0. 23
45	HEMBA1004433	F-HEMBA1004433	0.51
	HEMBA1004577	F-HEMBA1004577	0. 51
	HEMBA1004730	F-HEMBA1004730	0. 55
	HEMBA1004778	F-HEMBA1004778	0.47
50	HEMBA1004803	F-HEMBA1004803	0. 36
	HEMBA1004807	F-HEMBA1004807	0. 48
	HEMBA1004880	F-HEMBA1004880	0. 43
	HEMBA1004900	F-HEMBA1004900	0.40
	HEMBA1004909	F-HEMBA1004909	0. 24
55	HEMBA1004983	F-HEMBA1004983	0. 35

	HEMBA1005123	F-HEMBA1005123	0. 35
	HEMBA1005232	F-HEMBA1005232	0.09
	HEMBA1005241	F-HEMBA1005241	0. 34
5	HEMBA1005311	F-HEMBA1005311_	
	HEMBA1005318	F-HEMBA1005318	0. 59
	HEMBA1005353	F-HEMBA1005353	0. 34
	HEMBA1005374	F-HEMBA1005374	0. 38
10	HEMBA1005447	F-HEMBA1005447	0. 34
	HEMBA1005508	F-HEMBA1005508	0. 14
	HEMBA1005577	F-HEMBA1005577	0. 22
	HEMBA1005588	F-HEMBA1005588	0. 54
15	HEMBA1005593	F-HEMBA1005593	0. 44
· ·	HEMBA1005606	F-HEMBA1005606	0. 38
	HEMBA1005679	F-HEMBA1005679,	
	HEMBA1005894	F-HEMBA1005894	0. 46
20	HEMBA1005911	F-HEMBA1005911	0. 56
20	HEMBA1006036	F-HEMBA1006036	0. 87
	HEMBA1006124	F-HEMBA1006124	0. 35
	HEMBA1006253	F-HEMBA1006253	0. 35
0.5	HEMBA1006259	F-HEMBA1006259	0. 54
25	HEMBA1006364	F-HEMBA1006364	0. 55
	HEMBA1006380	F-HEMBA1006380	0. 48
	HEMBA1006426	F-HEMBA1006426	0. 54
	HEMBA1006461	F-HEMBA1006461	0. 20
30	HEMBA1006562	F-HEMBA1006562	0. 72
	HEMBA1006597	F-HEMBA1006597	0. 64
	HEMBA1006639	F-HEMBA1006639	0. 55
	HEMBA1006653	F-HEMBA1006653	0. 34
35	HEMBA1006695	F-HEMBA1006695	0.16
	HEMBA1006696	F-HEMBA1006696	0.37
	HEMBA1006744	F-HEMBA1006744	0. 33
	HEMBA1006824	F-HEMBA1006824	0. 37
40	HEMBA1006949	F-HEMBA1006949	0. 78
	HEMBA1007078	F-HEMBA1007078	0. 44
	HEMBA1007129	F-HEMBA1007129	0. 55
	HEMBA1007147	F-HEMBA1007147	0.41
45	HEMBA1007206	F-HEMBA1007206	0. 46
	HEMBA1007279	F-HEMBA1007279	0. 45
	HEMBA1007327	F-HEMBA1007327	0. 71
	HEMBB1000005	F-HEMBB1000005	0. 35
50	HEMBB1000054	F-HEMBB1000054	0. 18
	HEMBB1000055	F-HEMBB1000055	0. 87
	HEMBB1000144	F-HEMBB1000144	0. 47
	HEMBB1000258	F-HEMBB1000258	0. 87
55	HEMBB1000318	F-HEMBB1000318	0. 34
	HEMBB1000335	F-HEMBB1000335	0. 48

	HEMBB1000336	F-HEMBB1000336	0. 35
	HEMBB1000354	F-HEMBB1000354	0. 47
	HEMBB1000374	F-HEMBB1000374	0.69
5	HEMBB1000402	F-HEMBB1000402	0. 54
	HEMBB1000404	F-HEMBB1000404	0. 47
	HEMBB1000480	F-HEMBB1000480	0. 35
	HEMBB1000493	F-HEMBB1000493	0. 33
10	HEMBB1000554	F-HEMBB1000554	0.46
	HEMBB1000573	F-HEMBB1000573	0.34
	HEMBB1000649	F-HEMBB1000649	0. 41
	HEMBB1000652	F-HEMBB1000652	0. 46
15	HEMBB1000709	F-HEMBB1000709	0. 56
15	HEMBB1000749	F-HEMBB1000749	0. 36
	HEMBB1000790	F-HEMBB1000790,	
	HEMBB1000827	F-HEMBB1000827	0.31
	HEMBB1000831	F-HEMBB1000831	0. 42
20	HEMBB1000893	F-HEMBB1000893	0. 36
	HEMBB1001004	F-HEMBB1001004	0. 43
	HEMBB1001008	F-HEMBB1001008	0. 44
	HEMBB1001047	F-HEMBB1001047	0. 47
25	HEMBB1001119	F-HEMBB1001119	0. 25
	HEMBB1001142	F-HEMBB1001142	0. 15
	HEMBB1001315	F-HEMBB1001315	0. 72
	HEMBB1001317	F-HEMBB1001317	0. 72
30	HEMBB1001317	F-HEMBB1001326	0. 44
	HEMBB1001327	F-HEMBB1001337	0. 18
	HEMBB1001367	F-HEMBB1001367	0. 18
	HEMBB1001307	F-HEMBB1001424	0. 34
35	HEMBB1001436	F-HEMBB1001436	0.34
	HEMBB1001458	F-HEMBB1001458	0. 47
	HEMBB1001535	F-HEMBB1001535	0. 34
	HEMBB1001536	F-HEMBB1001536	0. 22
40	HEMBB1001565	F-HEMB81001565	0. 35
40	HEMBB1001747	F-HEMBB1001747	0.40
	HEMBB1001749		
	HEMBB1001797	F-HEMBB1001797	0. 79
	HEMBB1001836	F-HEMBB1001836	0. 52
45	HEMBB1001863	F-HEMBB1001863	0. 32
	HEMBB1001868	F-HEMBB1001868	0. 33
	HEMBB1001875	F-HEMBB1001875	0. 46
	HEMBB1001911	F-HEMBB1001911	0. 46
50	HEMBB1001911	F-HEMBB1001911	0. 38
	HEMBB1001925	F-HEMBB1001925	0. 33
	HEMBB1001944	F-HEMBB1001944	0. 35
	HEMBB1001983	F-HEMBB1001983	0.35
55	HEMBB1001983	F-HEMBB1001996	0.32
	115000 100 1330	L TIEMBOIOUIS90	U. 39

	HEMBB1001997	F-HEMBB1001997	0. 36
	HEMBB1002045	F-HEMBB1002045	0. 20
	HEMBB1002092	F-HEMBB1002092	0.31
5	HEMBB1002247	F-HEMBB1002247	
	HEMBB1002266	F-HEMBB1002266	0.34
	HEMBB1002387	F-HEMBB1002387	0. 32
	HEMBB1002425	F-HEMBB1002425	0. 31
10	HEMBB1002458	F-HEMBB1002458	0. 44
	HEMBB1002522	F-HEMBB1002522	0. 56
	HEMBB1002534	F-HEMBB1002534	0.43
	HEMBB1002579	F-HEMBB1002579	0, 21
15	HEMBB1002582	F-HEMBB1002582	0, 48
	HEMBB1002596	F-HEMBB1002596	0. 42
	HEMBB1002617	F-HEMBB1002617,	
	HEMBB1002702	F-HEMBB1002702	0. 31
20	MAMMA1000043	F-MAMMA1000043	0.45
20	MAMMA1000092	F-MAMMA1000092	0.39
	MAMMA1000129	F-MAMMA1000129	0.47
	MAMMA1000175	F-MAMMA1000175	0.18
	MAMMA1000198	F-MAMMA1000198	0.46
25	MAMMA1000221	F-MAMMA1000221	0.43
	MAMMA1000307	F-MAMMA1000307	0.35
	MAMMA1000331	F-MAMMA1000331	0.41
	MAMMA1000356	F-MAMMA1000356	0.41
30	MAMMA1000360	F-MAMMA1000360	0. 33
	MAMMA1000402	F-MAMMA1000402	0. 35
	MAMMA1000414	F-MAMMA1000414	0. 32
	MAMMA1000444	F-MAMMA1000444	0. 13
35	MAMMA1000500	F-MAMMA1000500	0. 36
	MAMMA1000522	F-MAMMA1000522	0.46
	MAMMA1000576	F-MAMMA1000576	0. 58
	MAMMA1000594	F-MAMMA1000594	0.69
40	MAMMA1000597	F-MAMMA1000597	0. 35
	MAMMA1000720	F-MAMMA1000720	0.43
	MAMMA1000761		0. 13
	MAMMA1000775	F-MAMMA1000775	0. 35
45	MAMMA1000778	F-MAMMA1000778	0. 54
	MAMMA1000798	F-MAMMA1000798	0. 40
	MAMMA1000862	F-MAMMA1000862	0. 87
	MAMMA1000876	F-MAMMA1000876	0. 47
50	MAMMA1000883	F-MAMMA1000883	0. 41
	MAMMA1000931	F-MAMMA1000931	0. 45
	MAMMA1000940	F-MAMMA1000940	0. 35
	MAMMA1000941	F-MAMMA1000941	0. 35
55	MAMMA1000943	F-MAMMA1000943	0. 23
	MAMMA1000975	F-MAMMA1000975	0.32

	MAMMA1001038	F-MAMMA1001038	0. 41
	MAMMA1001162	F-MAMMA1001162	0.18
	MAMMA1001186	F-MAMMA1001186	0.49
5	MAMMA1001220	F-MAMMA1001220	0. 43
	MAMMA1001256	F-MAMMA1001256	0. 35
	MAMMA1001274	F-MAMMA1001274	0. 57
	MAMMA1001341	F-MAMMA1001341	0. 45
10	MAMMA1001397	F-MAMMA1001397	0. 64
	MAMMA1001420	F-MAMMA1001420	0. 41
	MAMMA1001547	F-MAMMA1001547	0.39
	MAMMA1001590	F-MAMMA1001590	0. 46
	MAMMA1001670	F-MAMMA1001670	0. 46
15			
	MAMMA1001679	F-MAMMA1001679	0.46
	MAMMA1001711	F-MAMMA1001711,	
	MAMMA1001745	F-MAMMA1001745	0.53
20	MAMMA1001760	F-MAMMA1001760	0.58
	MAMMA1001769	F-MAMMA1001769	0. 33
	MAMMA1001815	F-MAMMA1001815	0. 43
	MAMMA1001820	F-MAMMA1001820	0.19
25	MAMMA1001907	F-MAMMA1001907	0. 38
	MAMMA1002056	F-MAMMA1002056	0. 35
	MAMMA1002078	F-MAMMA1002078	0.34
	MAMMA1002093	F-MAMMA1002093	0. 55
30	MAMMA1002125	F-MAMMA1002125	0. 39
30	MAMMA1002132	F-MAMMA1002132	0. 57
	MAMMA1002145	F-MAMMA1002145	0.78
	MAMMA1002250	F-MAMMA1002250	0.72
	MAMMA1002311	F-MAMMA1002311	0.58
35	MAMMA1002360	F-MAMMA1002360	0.19
	MAMMA1002411	F-MAMMA1002411	0.40
	MAMMA1002498	F-MAMMA1002498	0. 37
	MAMMA1002571	F-MAMMA1002571	0. 32
40	MAMMA1002701	F-MAMMA1002701	0.55
	MAMMA1002727	F-MAMMA1002727	0.34
	MAMMA1002728	F-MAMMA1002728	0. 55
	MAMMA1002746	F-MAMMA1002746	0.45
45	MAMMA1002764	F-MAMMA1002764	0.44
	MAMMA1002765	F-MAMMA1002765	0.40
	MAMMA1002820	F-MAMMA1002820	0.32
	MAMMA1002830	F-MAMMA1002830	0.36
50	MAMMA1002909	F-MAMMA1002909	0.64
JU	MAMMA1002941	F-MAMMA1002941	0.42
	MAMMA1002972	F-MAMMA1002972	0.08
	MAMMA1002973	F-MAMMA1002973	0.35
	MAMMA1003004	F-MAMMA1003004	0. 42
55	MAMMA1003007	F-MAMMA1003007	0.41

	MAMMA1003039	F-MAMMA1003039	0. 46
	MAMMA1003089		0. 58
	NT2RM4000086		0. 36
5	NT2RM4000265		
	NT2RM4000366	~ ~	0. 21
	NT2RM4000414		0. 35
	NT2RM4000425		0. 22
10	NT2RM4000779		0. 43
	NT2RM4000855		0. 52
	NT2RM4001160		0. 36
	NT2RM4001313		0. 38
15	NT2RM4001437		0. 53
	NT2RM4001754		0.35
	NT2RM4001856		
	NT2RM4001953		0. 59
20	NT2RM4001984		0. 35
20	NT2RM4002390		0. 24
	NT2RP2000077		0. 35
	NT2RP2000108	F-NT2RP2000108	0.08
25	NT2RP2000183	F-NT2RP2000183	0. 43
25	NT2RP2000257	F-NT2RP2000257	0.07
	NT2RP2000289	F-NT2RP2000289	0.35
	NT2RP2000420	F-NT2RP2000420	0.48
	NT2RP2000678	F-NT2RP2000678	0. 55
30	NT2RP2000715	F-NT2RP2000715	0. 37
	NT2RP2000842	F-NT2RP2000842	0.40
	NT2RP2000970	F-NT2RP2000970	0.46
	NT2RP2001149	F-NT2RP2001149	0.31
35	NT2RP2001226	F-NT2RP2001226	0. 44
	NT2RP2001295	F-NT2RP2001295	0.40
	NT2RP2001347	F-NT2RP2001347	0.41
	NT2RP2001467	F-NT2RP2001467	0.44
40	NT2RP2001506	F-NT2RP2001506	0. 17
	NT2RP2001569	F-NT2RP2001569	0. 31
	NT2RP2001663	F-NT2RP2001663	0. 37
		F-NT2RP2001936	0. 33
45	NT2RP2002041		0. 31
	NT2RP2002047		0. 15
	NT2RP2002066	F-NT2RP2002066	0.11
	NT2RP2002172	F-NT2RP2002172	0. 32
50	NT2RP2002219	F-NT2RP2002219	0. 54
	NT2RP2002316	F-NT2RP2002316	0.39
	NT2RP2002475	F-NT2RP2002475	0.09
	NT2RP2002546	F-NT2RP2002546	0. 33
55	NT2RP2002591	F-NT2RP2002591	0. 38
-	NT2RP2002643	F-NT2RP2002643	0. <b>5</b> 5

	NT2RP2002741	F-NT2RP2002741	0. 73
	NT2RP2002750	F-NT2RP2002741	0. 73
	NT2RP2002778		0.40
5	NT2RP2002778		
	NT2RP2002857		-
		F-NT2RP2003000	0.56
	NT2RP2003073	F-NT2RP2003073	0.36
10	NT2RP2003237	F-NT2RP2003237	0.69
70	NT2RP2003394	F-NT2RP2003394	0.34
	NT2RP2003517	F-NT2RP2003517	0.44
	NT2RP2003668	F-NT2RP2003668	0. 46
	NT2RP2003988	F-NT2RP2003988	0.55
15	NT2RP2004232		0.34
	NT2RP2004400		0.18
	NT2RP2004512	,	
	NT2RP2004523		
20	NT2RP2004587		0.09
	NT2RP2004736	F-NT2RP2004736	0. 70
	NT2RP2004767	F-NT2RP2004767	0. 54
	NT2RP2004775	F-NT2RP2004775	0. 53
25	NT2RP2004961	F-NT2RP2004961	0.49
	NT2RP2004962	F-NT2RP2004962	0.87
	NT2RP2004982	F-NT2RP2004982	0. 32
	NT2RP2005289	F-NT2RP2005289	0. 24
30	NT2RP2005407	F-NT2RP2005407	0. 33
	NT2RP2005531	F-NT2RP2005531	0. 20
	NT2RP2005694	F-NT2RP2005694	0.21
	NT2RP2005726	F-NT2RP2005726	0.42
	NT2RP2005942	F-NT2RP2005942	0.10
35	NT2RP2006258	F-NT2RP2006258	0. 37
	NT2RP2006261	F-NT2RP2006261	0. 42
	NT2RP2006454		0.46
	NT2RP2006554		0. 14
40	NT2RP3000055	F-NT2RP3000055	0.31
	NT2RP3000233	F-NT2RP3000233	0.54
	NT2RP3000341	F-NT2RP3000341	0.43
	NT2RP3000418	F-NT2RP3000418	0.53
45	NT2RP3000451	F-NT2RP3000451	0.46
	NT2RP3000561	F-NT2RP3000561	0. 47
	NT2RP3000582	F-NT2RP3000582	0.44
	NT2RP3001007	F-NT2RP3001007	0.26
50	NT2RP3001281	F-NT2RP3001281	0.36
	NT2RP3001318	F-NT2RP3001318	0.21
	NT2RP3001339	F-NT2RP3001339	0. 58
	NT2RP3001340	F-NT2RP3001340	0.41
55	NT2RP3001383	F-NT2RP3001383	0.36
55	NT2RP3001432	F-NT2RP3001432	0.34

	NT2RP3001580	F-NT2RP3001580	0.45
	NT2RP3001589	F-NT2RP3001589	0.48
	NT2RP3001898	F-NT2RP3001898	0.19
5	NT2RP3002004	F-NT2RP3002004	0.36
	NT2RP3002173	F-NT2RP3002173	0.39
	NT2RP3003133		0. 35
	NT2RP3003264		0. 13
10	NT2RP3003346		0. 72
	NT2RP3003403		0. 45
	NT2RP3003433		0. 24
	NT2RP3003576		0. 35
15	NT2RP3003625		0. 43
15	NT2RP3003665		0. 32
	NT2RP3003800		
	NT2RP3003800 NT2RP3003828		0. 52
	NT2RP3003828 NT2RP3003842		0. 32
20	NT2RF3004070		0. 23
	NT2RF3004470		0. 32
	NT2RP4000023		0. 46
	NT2RP4000025		0. 56
25	NT2RP4000102		0.38
	NT2RP4000102 NT2RP4000167		
	NT2RP4000167		0.42
			0.33
30	NT2RP4000218		0.37
	NT2RP4000424		0.48
	NT2RP4000915		0.37
	NT2RP4002075		0.43
25	0VARC1000017		0.14
35	0VARC1000068		0.17
	0VARC1000085	F-0VARC1000085	0. 46
	0VARC1000092		0. 58
	0VARC1000145	F-0VARC1000145	0.53
40	0VARC1000414	F-0VARC1000414	0.40
	0VARC1000486	F-0VARC1000486	0. 15
	0VARC1000496	1 01/4(01000400	
	0VARC1000526		0. 33
45	0VARC1000948		0.71
	0VARC1001011	F-0VARC1001011	0. 42
	0VARC1001240	F-0VARC1001240	0. 14
	0VARC1001600	F-0VARC1001600	0. 37
50	0VARC1001805	F-0VARC1001805	0.51
	OVARC1001813	F-0VARC1001813	0. 58
	0VARC1001846	F-0VARC1001846	0. 59
	PLACE1000540	F-PLACE1000540	0. 35
	PLACE1000599	F-PLACE1000599	0. 37
55	PLACE1001088	F-PLACE1001088	0.36

	PLACE1001323	F-PLACE1001323	0. 21
	PLACE1001377		0. 41
	PLACE1001440	F-PLACE1001440	0. 45
5	PLACE1001517	F-PLACE1001517	0, 41
	PLACE1001672	F-PLACE1001672	0. 35
	PLACE1001705	F-PLACE1001705	0.11
	PLACE1001756	F-PLACE1001756	0. 67
10	PLACE1002157	F-PLACE1002157	0. 68
	PLACE1002205	F-PLACE1002105	0. 55
	PLACE1002227		0. 33
	PLACE1002227 PLACE1002259	. , _,,,_,,	0. 14
15	PLACE1002319	F-PLACE1002319	0. 29
	PLACE1002399		0.31
	PLACE1002477	F-PLACE1002477	
	PLACE1002500	F-PLACE1002500	0. 26
20	PLACE1002583	F-PLACE1002583	0.49
	PLACE1002604	F-PLACE1002604	0. 25
	PLACE1002772		0. 13
	PLACE1002853	F-PLACE1002853	0. 46
25	PLACE1002968	F-PLACE1002968	0. 33
	PLACE1003238	F-PLACE1003238	0. 46
	PLACE1003420	F-PLACE1003420	0. 42
	PLACE1003478	F-PLACE1003478	0. 20
20	PLACE1003566	F-PLACE1003566	0. 35
30	PLACE1003593	F-PLACE1003593	0. 42
	PLACE1003618	F-PLACE1003618	0.36
	PLACE1004274		0. 41
	PLACE1004681	F-PLACE1004681	0. 25
35	PLACE1004716	F-PLACE1004716	0. 49
	PLACE1004773	F-PLACE1004773	0. 31
	PLACE1004815	F-PLACE1004815	0. 72
	PLACE1004836	F-PLACE1004836	0. 39
40	PLACE1004913	F-PLACE1004913	0. 44
	PLACE1004979	F-PLACE1004979	0. 44
	PLACE1005052	F-PLACE1005052	0. 63
	PLACE1005086	F-PLACE1005086	0. 57
45	PLACE1005108	F-PLACE1005108	0. 17
	PLACE1005128	F-PLACE1005128	0.40
	PLACE1005176	F-PLACE1005176	0. 48
	PLACE1005467	F-PLACE1005467	0.37
	PLACE1005639	F-PLACE1005639	0. 33
50	PLACE1005850	F-PLACE1005850	0. 54
	PLACE1005932	F-PLACE1005932	0.20
	PLACE1006003	F-PLACE1006003	0.71
	PLACE1006017	F-PLACE1006017	0. 42
55	PLACE1006288	F-PLACE1006288	0.34

	PLACE1006318	F-PLACE1006318	0. 22
	PLACE1006368	F-PLACE1006368	0.10
	PLACE1006371	F-PLACE1006371	0.43
5	PLACE1006506	F-PLACE1006506	0. 28
	PLACE1006629	F-PLACE1006629	0.44
	PLACE1006795	F-PLACE1006795	0. 35
	PLACE1006904	F-PLACE1006904	0. 20
10	PLACE1000904	F-PLACE1007478	0. 70
70	PLACE1007478		
		F~PLACE1007557	0.17
	PLACE1007743	F-PLACE1007743	0. 28
	PLACE1007829	F-PLACE1007829	0.16
15	PLACE1007877	F-PLACE1007877	0.16
	PLACE1008181	F-PLACE1008181	0. 39
	PLACE1008330	F-PLACE1008330,	
	PLACE1008584	F-PLACE1008584	0, 61
20	PLACE1008630	F-PLACE1008630	0.09
	PLACE1008715	F-PLACE1008715	0. 36
	PLACE1008851	F-PLACE1008851	0. 39
	PLACE1008941	F-PLACE1008941	0. 42
25	PLACE1009039	F-PLACE1009039	0. 43
25	PLACE1009048	F-PLACE1009048	0. 24
	PLACE1009493	F-PLACE1009493	0. 35
	PLACE1009539	F-PLACE1009539	0.31
	PLACE1009637	F-PLACE1009637	0. 58
30	PLACE1009925	F-PLACE1009925	0.17
	PLACE1009947	F-PLACE1009947	0.42
	PLACE1010231	F-PLACE1010231	0.36
	PLACE1010562	F-PLACE1010562	0. 63
35	PLACE1010579	F-PLACE1010579	0. 42
	PLACE1010739	F-PLACE1010739	0.49
	PLACE1010802	F-PLACE1010802	0.41
	PLACE1010896	F-PLACE1010896	0.47
40	PLACE1011032	F-PLACE1011032	0.46
40	PLACE1011109	F-PLACE1011109	0.19
	PLACE1011185	F-PLACE1011185	0.66
	PLACE1011452	F-PLACE1011452	0.72
	PLACE1011465	F-PLACE1011465	0. 51
45	PLACE1011492	F-PLACE1011492	0. 19
	PLACE1011520	F-PLACE1011520	0. 54
	PLACE1011567	F-PLACE1011567	0.51
	PLACE1011643	F-PLACE1011643	0. 21
50	PLACE1011643	F-PLACE1011719	0. 21
	PLACE10117/19 PLACE10117/49		
		F-PLACE1011749	0. 24
	PLACE2000011	F-PLACE2000011	0. 48
	PLACE2000017	F-PLACE2000017	0. 34
55	PLACE2000061	F-PLACE2000061	0. 34

	PLACE2000187	F-PLACE2000187	0, 40
	PLACE2000216	F-PLACE2000216	0. 32
	PLACE2000335	F-PLACE2000335	0. 78
5	PLACE2000347	F-PLACE2000347	0. 36
	PLACE2000366	F-PLACE2000366	0. 33
	PLACE2000394	F-PLACE2000394	0. 54
	PLACE2000398	F-PLACE2000398	0. 44
10	PLACE2000425	F-PLACE2000425	0. 47
	PLACE2000450	F-PLACE2000450	0. 53
	PLACE2000477	F-PLACE2000477	0.36
	PLACE3000119	F-PLACE3000119	0.40
15	PLACE3000207	F-PLACE3000207	0. 32
75	PLACE3000230	F-PLACE3000230	0. 51
	PLACE3000271	F-PLACE3000271,	
	PLACE3000373	F-PLACE3000373	0. 37
	PLACE3000399	F-PLACE3000399	0. 42
20	PLACE3000401	F-PLACE3000401	0. 37
	PLACE3000406	F-PLACE3000406	0. 46
	PLACE4000247	F-PLACE4000247	0. 64
	PLACE4000320	F-PLACE4000320	0. 36
25	PLACE4000344	F-PLACE4000344	0. 31
	PLACE4000367	F-PLACE4000367	0. 32
	PLACE4000401	F-PLACE4000401	0. 54
	PLACE4000548	F-PLACE4000548	0. 26
30	THYR01000111	F-THYR01000111	0. 53
	THYR01000129	F-THYR01000129	0. 45
	THYRO1000187	F-THYR01000187	0. 46
	THYR01000279	F-THYR01000279	0.09
35	THYR01000484	F-THYR01000484	0.36
	THYR01000596	F-THYR01000596	0. 47
	THYR01000625	F-THYR01000625	0. 59
	THYR01000793	F-THYR01000793	0. 17
40	THYR01000815	F-THYR01000815	0. 58
	THYR01000865	F-THYR01000865	0. 55
	THYR01001003	F-THYR01001003	0.34
	THYR01001031	F-THYR01001031	0. 43
45	THYR01001133	F-THYR01001133	0. 52
	THYR01001321	F-THYR01001321	0.37
	THYR01001401	F-THYR01001401	0. 36
	THYR01001426	F-THYR01001426	0.32
50	THYR01001434	F-THYR01001434	0. 39
	THYR01001559	F-THYR01001559	0. 45
	THYR01001570	F-THYR01001570	0. 35
	THYR01001706 THYR01001746	F-THYR01001706	0. 45
er.		F-THYR01001746	0. 43
55	THYR01001772	F-THYR01001772	0. 37

THYR01001907	F-THYR01001907	0. 33
Y79AA1000033	F-Y79AA1000033	0.53
Y79AA1000346	F-Y79AA1000346	0.53
Y79AA1000410	F-Y79AA1000410	0. 13
Y79AA1000805	F-Y79AA1000805	0.32
Y79AA1001692	F-Y79AA1001692	0. 45
Y79AA1002103	F-Y79AA1002103	0. 15
Y79AA1002220	F-Y79AA1002220	0.49
		VV

#### 15 EXAMPLE 13

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Full-length sequence analysis and homology search

[0192] Full-length sequence was determined for each selected cDNA clones. The nucleotide sequence determination was performed mainly by the dye-terminator method using custom synthesized DNA primers according to the primer walking procedure (custom synthesized DNA primers were used for sequencing; sequencing reaction was performed with DNA sequencing reagent supplied by PE Biosystems according to the supplier's manual; and the samples were analyzed in an automatic sequencer made by the same supplier). Sequence determination of some clones was carried out in the same manner but using a Licor DNA sequencer. Overlapping partial nucleotide sequences, which were obtained by the above-described method, were assembled together to determine a full-length nucleotide sequence. Amino acid sequences were then deduced from the determined full-length nucleotide sequences. However, amino acid sequence is not shown for a clone of which coding region was hard to be deduced or of which amino acid sequence has less than 100 amino acid residues. SEQ ID NOs corresponding to the respective clones are indicated in Table 350 and Table 351.

[0193] GenBank, Swiss-Prot and UniGene were searched for the determined nucleotide sequences by BLAST analysis. Matching data of cDNA clone which exhibits higher homology and of which functions are easily predicted based on the nucleotide sequences and the deduced amino acid sequences are selected from the BLAST analysis matching data with P value of 10<sup>-4</sup> or less. The matching data selected are listed herein. The results of homology search 6, 12, 13, and 14 are indicated in the last part of this specification. However, there are some clones that did not match the criteria for judgment and such matching data of BLAST analysis are not shown herein.

#### **EXAMPLE 14**

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Novel full-length cDNA clone obtained from a cDNA library prepared by oligo-capping method

[0194] A cDNA clone, NT2RP4002298, was obtained from a cDNA library, NT2RP4 (see Example 1), prepared by oligo-capping method. Analysis of the entire nucleotide sequence of the clone has revealed that the clone encodes a novel protein consisting of 775 amino acids. The ATGpr1 score at the initiation codon of the amino acid sequence is 0.16, and therefore, the fullness ratio is low. However, the sequence can be full-length.

[0195] The full-length nucleotide sequence of NT2RP4002298 is shown in SEQ ID NO: 12370, and the deduced amino acid sequence encoded by the clone NT2RP4002298 is shown in SEQ ID NO: 12371.

#### **EXAMPLE 15**

Gene expression analysis with hybridization using high density DNA filter

[0196] Nylon membrane for DNA spotting was prepared according to the following procedure. E. coli was cultured in each well of a 96-well plate (in a LB medium at  $37^{\circ}$ C for 16 hours). A small aliquot of each culture was suspended in 10  $\mu$ l of sterile water in a well of a 96-well plate. The plate was heated at  $100^{\circ}$ C for 10 minutes. Then the boiled samples were analyzed by PCR reaction. PCR was performed in a 20  $\mu$ l solution by using TaKaRa PCR Amplification Kit (Takara) according to the supplier's protocol. Primers used for the amplification of an insert cDNA in a plasmid were a pair of sequencing primers, ME761FW (5' tacggaagtgttacttctgc 3' / SEQ ID NO: 13290) and ME1250RV (5' tgtgggaggttttttctcta 3' / SEQ ID NO: 13291), or a pair of primers, M13M4 (5' gttttcccagtcacgac 3' / SEQ ID NO: 13292)

and M13RV (5' caggaaacagctatgac 3' / SEQ ID NO: 13293). PCR reaction was performed in a thermal cycler, GeneAmp System 9600 (PE Biosystems). The cycling profile consisted of pre-heat at 95°C for 5 mimutes; 10 cycles of denaturation at 95°C for 10 seconds, and annealing/extension at 68°C for 1 minute; 20 cycles of denaturation at 98°C for 20 seconds and annealing/extension at 60°C for 3 minutes; and final extension at 72°C for 10 minutes. After the PCR reaction, the 20 µl reaction solution was loaded onto a 1 % agarose gel and fractionated by electrophoresis. DNA on the gel was stained with ethidium bromide to confirm the amplification of cDNA. When cDNAs were barely amplified by PCR, plasmids containing the corresponding insert cDNAs were prepared by the alkali-extraction method (J. Sambrook, E. F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

[0197] Preparation of DNA array was carried out by the following procedure. An Aliquot of a DNA solution was added in each well of a 384-well plate. DNA was spotted onto a nylon membrane (Boehringer) by using a 384-pin tool of Biomek 2000 Laboratory Automation Sysytem (Beckman-Coulter). Specifically, the 384-well plate containing the DNA was placed under the 384-pin tool. The independent 384 needles were simultaneously dipped into the DNA solution for DNA deposition. The needles were gently pressed onto a nylon membrane and the DNA deposited at the tips of needles was spotted onto the membrane. Denaturation of the spotted DNA and immobilization of the DNA on the nylon membrane were carried out according to usual manners (J. Sambrook, E.F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press. 1989).

[0198] Hybridization probe used was radioisotope-labeled 1st strand cDNA. The 1st strand cDNA synthesis was performed by using Thermoscript<sup>(TM)</sup> RT-PCR System (GIBCO). Specifically, the 1st strand cDNA was synthesized by using 1.5  $\mu$ g mRNAs from various human tissues (Clontech), 1  $\mu$ l aliquots of 50  $\mu$ M Oligo(dT)20 and 50  $\mu$ Ci [ $\alpha$  <sup>33</sup>P] dATP according to an attached protocol. Probe purification was carried out by using ProbeQuant<sup>(TM)</sup> G-50 micro column (Amersham-Pharmacia Biotech) according to an attached protocol. In the next step, 2 units of E. coli RNasH were added to the reaction mixture. The mixture was incubated at room temperature for 10 minutes and then 100  $\mu$ g of human COT-1 DNA (GIBCO) was added thereto. The mixture was incubated at 97°C for 10 minutes and then was allowed to stand on ice to give hybridization probe.

[0199] Hybridization of the radioisotope-labeled probe to the DNA array was performed in a usual manner (J. Sambrook, E.F., Fritsh, & T. Maniatis, Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989). The membrane was washed as follows: the nylon membrane was incubated in Washing solution 1 (2 × SSC, 1% SDS) at room temperature (about 26°C) for 20 minutes and this washing was repeated 3 times; then the membrane was washed 3 times by incubating it in Washing solution 2 (0.1 × SSC, 1% SDS) at 65°C for 20 minutes. Autoradiography was performed by using an image plate for BAS2000 (Fuji Photo Film Co., Ltd.). Specifically, the nylon membrane with probe hybridized thereon was wrapped with a piece of Saran Wrap and brought into tight contact with the image plate on the light-sensitive surface. The membrane with the image plate was placed in an imaging cassette for radioisotope and allowed to stand in dark place for 4 hours. The radioactivity recorded on the image plate was analyzed by using BAS2000 (Fuji Photo Film Co., Ltd.). The activity was subjected to electronic conversion and recorded as an image file of autoradiogram. The signal intensity of each DNA spot was analyzed by using Visage High Density Grid Analysis Systems (Genomic Solutions Inc.). The signal intensity was converted into numerical data. The data were taken by duplicated measurements. The reproducibility was assessed by comparing the signal intensities of the corresponding spots on the duplicated DNA filters that were hybridized to a single DNA probe (Figure 2). The ratio between the corresponding spots falls within a range of 2 or less in 95% of entire spots and the correlation coefficient is r=0.97. Thus the reproducibility is assumed to be satisfactory.

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[0200] The detection sensitivity in gene expression analysis was estimated by examining increases in the signal intensity of probe concentration-dependent spot in hybridization using a probe complementary to the DNA spotted on the nylon membrane. DNA used was PLACE1008092 (same as DNA deposited in GenBank under an Accession No. AF107253). The DNA array with DNA of PLACE1008092 was prepared according to the above-mentioned method. The probe used was prepared as follows: mRNA was synthesized in vitro from the clone, PLACE1008092. By using this mRNA as a template, radioisotope-labeled 1st strand cDNA was synthesized in the same manner as described above, and the cDNA was used as the probe. The cDNA PLACE1008092 was inserted into pBluescript SK(-), of which T7 promoter was ligated to the 5' end of the cDNA, to give a plasmid for in vitro synthesis of the mRNA from PLACE1008092. Specifically, the PLACE1008092 insert was cut out from pME18SFL3 carrying the cDNA at a DrallI site thereof by Xhol digestion. The resulting PLACE1008092 fragment was ligated to Xhol-predigested pBluescript SK (-) by using DNA ligation kit ver.2 (Takara). The in-vitro mRNA synthesis from PLACE1008092 inserted in pBluescript SK(-) was carried out by using Ampliscribe<sup>(TM)</sup> T7 high yield transcription kit (Epicentre technologies). Hybridization and the analysis of signal intensity of each DNA spot were conducted by the same methods as described above. When the probe concentration is  $1 \times 10^7$  µg/ml or less, there was no increase of signal intensity proportional to the probe concentration. Therefore it was assumed to be difficult to compare the signals with one another in the concentration range. Thus the spots with the intensity of 40 or less were indiscriminately taken as low-level signals (Figure 3). Within a concentration of the probe ranging from  $1 \times 10^7 \,\mu\text{g/ml}$  to 0.1  $\mu\text{g/ml}$ , the signal was found to increase in a probe

concentration-dependent manner. The detection limit was then assumed to be 1:100,000 in a ratio of mRNA expression level in a sample.

[0201] Tables 12-168 (also containing clones with no description in Examples) show the expression of each cDNA in human normal tissues (heart, lung, pituitary gland, thymus, brain, kidney, liver and spleen). The expression levels are indicated by numerical values of 0-10,000. Genes that were expressed in at least a single tissue are indicated below by the corresponding clone names:

	HEMBA1000012,	HEMBA1000020,	HEMBA1000030,	HEMBA1000042,	HEMBA1000046,
10	HEMBA1000076,	HEMBA1000111,	HEMBA1000129,	HEMBA1000150,	HEMBA1000156,
	HEMBA1000158,	HEMBA1000168,	HEMBA1000185,	HEMBA1000216,	HEMBA1000227,
	HEMBA1000231,	HEMBA1000243,	HEMBA1000244,	HEMBA1000280,	HEMBA1000282,
	HEMBA1000288,	HEMBA1000303,	HEMBA1000304,	HEMBA1000327,	HEMBA1000338,
15	HEMBA1000351,	HEMBA1000355,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000369,	HEMBA1000376,	HEMBA1000387,	HEMBA1000392,	HEMBA1000396,
	HEMBA1000411,	HEMBA1000422,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000469,	HEMBA1000488,	HEMBA1000491,	HEMBA1000501,	HEMBA1000505,
20	HEMBA1000508,	HEMBA1000519,	HEMBA1000523,	HEMBA1000534,	HEMBA1000540,
	HEMBA1000542,	HEMBA1000545,	HEMBA1000557,	HEMBA1000561,	HEMBA1000568,
	HEMBA1000569,	HEMBA1000575,	HEMBA1000588,	HEMBA1000591,	HEMBA1000604,

	-HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000657,	HEMBA1000673,
	HEMBA1000682,	HEMBA1000702,	HEMBA1000705,	HEMBA1000726,	HEMBA1000749,
	HEMBA1000752,	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,
5	HEMBA1000843,	HEMBA1000852,	HEMBA1000870,	HEMBA1000872,	HEMBA1000876,
	HEMBA1000908,	HEMBA1000910,	HEMBA1000918,	HEMBA1000919.	HEMBA1000934.
	HEMBA1000942,	HEMBA1000960,	HEMBA1000971,	HEMBA1000972,	HEMBA1000975,
	HEMBA1000986,	HEMBA1000991,	HEMBA1001007,	HEMBA1001008,	HEMBA1001017,
10	HEMBA1001020,	HEMBA1001051, HE			
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001121,
	HEMBA1001123,	HEMBA1001133,	HEMBA1001137,	HEMBA1001140,	HEMBA1001172,
	HEMBA1001197,	HEMBA1001208,	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,
15	HEMBA1001265,	HEMBA1001281,	HEMBA1001286,	HEMBA1001289,	HEMBA1001294,
	HEMBA1001299,	HEMBA1001302,	HEMBA1001303,	HEMBA1001310,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,,	HEMBA1001375,	HEMBA1001377,	HEMBA1001387,
	HEMBA1001388,	HEMBA1001398,	HEMBA1001405,	HEMBA1001413,	HEMBA1001415,
20	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
	HEMBA1001454,	HEMBA1001463,	HEMBA1001476,	HEMBA1001497,	HEMBA1001510-
	HEMBA1001515,	HEMBA1001517,	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,
	HEMBA1001566,	HEMBA1001569,	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,
25	HEMBA1001589,	HEMBA1001595,	HEMBA1001608,	HEMBA1001620,	HEMBA1001636;
23	HEMBA1001640,	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,	HEMBA1001661,
	HEMBA1001678,	HEMBA1001709,	HEMBA1001712,	HEMBA1001714,	HEMBA1001718,
	HEMBA1001731,	HEMBA1001734,	HEMBA1001745,	HEMBA1001761,	HEMBA1001781,
	HEMBA1001784,	HEMBA1001791,	HEMBA1001804,	HEMBA1001808,	HEMBA1001809,
30	HEMBA1001815,	HEMBA1001822,	HEMBA1001824,	HEMBA1001844,	HEMBA1001847,
	HEMBA1001864,	HEMBA1001866,	HEMBA1001869,	HEMBA1001888,	HEMBA1001896,
	HEMBA1001910,	HEMBA1001912,	HEMBA1001913,	HEMBA1001939,	HEMBA1001940,
	HEMBA1001964,	HEMBA1001967, HE	MBA1001987, HEMB	A1001991, HEMBA1	.002003,
35	HEMBA1002008,	HEMBA1002018,	HEMBA1002035,	HEMBA1002049,	HEMBA1002092,
	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119,	HEMBA1002125,
	HEMBA1002144,	HEMBA1002150,	HEMBA1002151,	HEMBA1002160,	HEMBA1002161,
	HEMBA1002162,	HEMBA1002166,	HEMBA1002177,	HEMBA1002185,	HEMBA1002189,
40	HEMBA1002191,	HEMBA1002226,	HEMBA1002229,	HEMBA1002237,	HEMBA1002241,
	HEMBA1002265,	HEMBA1002267,	HEMBA1002270,	HEMBA1002328,	HEMBA1002337,
	HEMBA1002341,	HEMBA1002363,	HEMBA1002381,	HEMBA1002389,	HEMBA1002417,
	HEMBA1002419,	HEMBA1002439,	HEMBA1002458,	HEMBA1002462,	HEMBA1002469,
45	HEMBA1002477,	HEMBA1002486,	HEMBA1002495,	HEMBA1002498,	HEMBA1002503,
	HEMBA1002508,	HEMBA1002513,	HEMBA1002515,	HEMBA1002538,	HEMBA1002542,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002561,	HEMBA1002569,	HEMBA1002590,
	HEMBA1002592,	HEMBA1002609,	HEMBA1002624,	HEMBA1002629,	HEMBA1002645,
50	HEMBA1002659,	HEMBA1002661,	HEMBA1002678,	HEMBA1002679,	HEMBA1002696,
	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,	HEMBA1002746,
	HEMBA1002748,	HEMBA1002750,	HEMBA1002768,	HEMBA1002777,	HEMBA1002779,
	HEMBA1002780,	HEMBA1002794,	HEMBA1002810,	HEMBA1002816,	HEMBA1002818,
55	HEMBA1002826,	HEMBA1002833,	HEMBA1002863,	HEMBA1002876,	HEMBA1002896,
33	HEMBA1002921,	HEMBA1002924,	HEMBA1002934,	HEMBA1002935,	HEMBA1002944,

	HEMBA1002968,	HEMBA1002970,	HEMBA1002971,	HEMBA1002973,	HEMBA1002997,
	HEMBA1003021,	HEMBA1003033, HE	MBA1003034, HEMB	A1003037, HEMBA1	003041,
	HEMBA1003046,	HEMBA1003067,	HEMBA1003071,	HEMBA1003078,	HEMBA1003079,
5	HEMBA1003083,	HEMBA1003086,	HEMBA1003098,	HEMBA1003129,	HEMBA1003133,
	HEMBA1003136,	HEMBA1003142,	HEMBA1003148,	HEMBA1003166,	HEMBA1003175,
	HEMBA1003179,	HEMBA1003199,	HEMBA1003202,	HEMBA1003204,	HEMBA1003212,
	HEMBA1003220,	HEMBA1003229,	HEMBA1003235,	HEMBA1003257,	HEMBA1003273,
10	HEMBA1003276,	HEMBA1003278,	HEMBA1003281,	HEMBA1003286,	HEMBA1003296,
	HEMBA1003314,	HEMBA1003322,	HEMBA1003327,	HEMBA1003328,	HEMBA1003330,
	HEMBA1003348,	HEMBA1003370,	HEMBA1003373,	HEMBA1003376,	HEMBA1003380,
	HEMBA1003384,	HEMBA1003403,	HEMBA1003408,	HEMBA1003418,	HEMBA1003447,
15	HEMBA1003461,	HEMBA1003463,	HEMBA1003480,	HEMBA1003528,	HEMBA1003531,
	HEMBA1003538,	HEMBA1003556,	HEMBA1003571,	HEMBA1003581,	HEMBA1003591,
	HEMBA1003595,	HEMBA1003597,	HEMBA1003598,	HEMBA1003615,	HEMBA1003617,
	HEMBA1003621,	HEMBA1003637,	HEMBA1003640,	HEMBA1003645,	HEMBA1003656,
20	HEMBA1003667,	HEMBA1003680,	HEMBA1003690,	HEMBA1003692,	HEMBA1003711,
	HEMBA1003714,	HEMBA1003715,	HEMBA1003720,	HEMBA1003725,	HEMBA1003729,
	HEMBA1003733,	HEMBA1003742,	HEMBA1003758,	HEMBA1003760,	HEMBA1003773,
	HEMBA1003803,	HEMBA1003804,	HEMBA1003805,	HEMBA1003827,	HEMBA1003836,
25	HEMBA1003838,	HEMBA1003864,	HEMBA1003879,	HEMBA1003880,	HEMBA1003885,
25	HEMBA1003893,	HEMBA1003902,	HEMBA1003926,	HEMBA1003937,	HEMBA1003942,
	HEMBA1003958,	HEMBA1003978, HEM	ABA1003987, HEMB	A1003989, HEMBA1	004011,
	HEMBA1004012,	HEMBA1004024,	HEMBA1004048,	HEMBA1004049,	HEMBA1004056,
	HEMBA1004074,	HEMBA1004086,	HEMBA1004097,	HEMBA1004111,	HEMBA1004131,
30	HEMBA1004132,	HEMBA1004133,	HEMBA1004138,	HEMBA1004146,	HEMBA1004164,
	HEMBA1004200,	HEMBA1004203,	HEMBA1004225,	HEMBA1004227,	HEMBA1004238,
	HEMBA1004246,	HEMBA1004267,	HEMBA1004274,	HEMBA1004275,	HEMBA1004289,
	HEMBA1004306,	HEMBA1004312,	HEMBA1004321,	HEMBA1004323,	HEMBA1004327,
35	HEMBA1004335,	HEMBA1004341,	HEMBA1004353,	HEMBA1004354,	HEMBA1004356,
	HEMBA1004396,	HEMBA1004405,	HEMBA1004408,	HEMBA1004429,	HEMBA1004433,
	HEMBA1004460,	HEMBA1004461,	HEMBA1004479,	HEMBA1004499,	HEMBA1004506,
	HEMBA1004507,	HEMBA1004509,	HEMBA1004534,	HEMBA1004538,	HEMBA1004542,
40	HEMBA1004554,	HEMBA1004560,	HEMBA1004577,	HEMBA1004586,	HEMBA1004596,
	HEMBA1004604,	HEMBA1004610,	HEMBA1004617,	HEMBA1004629,	HEMBA1004669,
•	HEMBA1004670,	HEMBA1004672,	HEMBA1004697,	HEMBA1004705,	HEMBA1004709,
	HEMBA1004711,	HEMBA1004725,	HEMBA1004736,	HEMBA1004748,	HEMBA1004751,
45	HEMBA1004752,	HEMBA1004753,	HEMBA1004756,	HEMBA1004758,	HEMBA1004763,
	HEMBA1004771,	HEMBA1004778,	HEMBA1004803,	HEMBA1004816,	HEMBA1004847,
	HEMBA1004850,	HEMBA1004863,	HEMBA1004864,	HEMBA1004865,	HEMBA1004880,
	HEMBA1004909,	HEMBA1004918,	HEMBA1004923,	HEMBA1004930,	HEMBA1004933,
50	HEMBA1004934,	HEMBA1004944,	HEMBA1004954,	HEMBA1004960,	HEMBA1004972,
	HEMBA1004980,	HEMBA1005008, HEM	IBA1005019, HEMB	A1005035, HEMBA1	
	HEMBA1005047,	HEMBA1005050,	HEMBA1005075,	HEMBA1005079,	HEMBA1005101,
	HEMBA1005113,	HEMBA1005123,	HEMBA1005133,	HEMBA1005149,	HEMBA1005152,
C E	HEMBA1005201,	HEMBA1005202,	HEMBA1005206,	HEMBA1005223,	HEMBA1005241,
55	HEMBA1005244,	HEMBA1005251,	HEMBA1005252,	HEMBA1005275,	HEMBA1005296,

	HEMBA1005304,	HEMBA10053-11,	HEMBA1005315,	HEMBA1005331,	HEMBA1005338,
	HEMBA1005353,	HEMBA1005359,	HEMBA1005367,	HEMBA1005374,	HEMBA1005382,
	HEMBA1005389,	HEMBA1005403,	HEMBA1005408,	HEMBA1005411,	HEMBA1005423,
5	HEMBA1005443,	HEMBA1005447,	HEMBA1005468,	HEMBA1005469,	HEMBA1005472,
	HEMBA1005474,	HEMBA1005475,	HEMBA1005500,	HEMBA1005511,	HEMBA1005513,
	HEMBA1005518,	HEMBA1005520,	HEMBA1005526,	HEMBA1005530,	HEMBA1005548,
	HEMBA1005552,	HEMBA1005558,	HEMBA1005568,	HEMBA1005570,	HEMBA1005576,
10	HEMBA1005577,	HEMBA1005581,	HEMBA1005583,	HEMBA1005588,	HEMBA1005593.
	HEMBA1005595,	HEMBA1005606,	HEMBA1005609,	HEMBA1005616,	HEMBA1005627,
	HEMBA1005632,	HEMBA1005634.	HEMBA1005670.	HEMBA1005679.	HEMBA1005680,
	HEMBA1005685,	HEMBA1005705,	HEMBA1005717,	HEMBA1005732,	HEMBA1005765,
15	HEMBA1005780,	HEMBA1005813,	HEMBA1005815,	HEMBA1005822,	HEMBA1005829,
, 0	HEMBA1005834,	HEMBA1005852,	HEMBA1005853,	HEMBA1005894,	HEMBA1005911,
	HEMBA1005921,	HEMBA1005931,	HEMBA1005934,	HEMBA1005962,	HEMBA1005990,
	HEMBA1005991,	HEMBA1005999,	HEMBA1006002,	HEMBA1006005,	HEMBA1006031.
20	HEMBA1006036,	HEMBA1006042, HE			
20	HEMBA1006100,	HEMBA1006108,	HEMBA1006121,	HEMBA1006124,	HEMBA1006138-
	HEMBA1006142,	HEMBA1006155,	HEMBA1006173,	HEMBA1006182,	HEMBA1006235,
	HEMBA1006248,	HEMBA1006252,	HEMBA1006253,	HEMBA1006259,	HEMBA1006278,
	HEMBA1006283,	HEMBA1006309,	HEMBA1006310,	HEMBA1006328,	HEMBA1006344;
25	HEMBA1006349,	HEMBA1006359,	HEMBA1006364,	HEMBA1006377,	HEMBA1006380,
	HEMBA1006381,	HEMBA1006398,	HEMBA1006416,	HEMBA1006419,	HEMBA1006421,
	HEMBA1006426,	HEMBA1006438,	HEMBA1006445,	HEMBA1006461,	HEMBA1006483,
	HEMBA1006485,	HEMBA1006486,	HEMBA1006497,	HEMBA1006502,	HEMBA1006546,
30	HEMBA1006562,	HEMBA1006569,	HEMBA1006595,	HEMBA1006597,	HEMBA1006612,
	HEMBA1006617,	HEMBA1006624,	HEMBA1006631,	HEMBA1006635,	HEMBA1006639,
	HEMBA1006643,	HEMBA1006648,	HEMBA1006652,	HEMBA1006653,	HEMBA1006659,
	HEMBA1006674,	HEMBA1006676,	HEMBA1006695,	HEMBA1006696,	HEMBA1006708,
35	HEMBA1006709,	HEMBA1006717,	HEMBA1006737,	HEMBA1006744,	HEMBA1006754,
	HEMBA1006758,	HEMBA1006779,	HEMBA1006780,	HEMBA1006795,	HEMBA1006796,
	HEMBA1006807,	HEMBA1006821,	HEMBA1006824,	HEMBA1006832,	HEMBA1006849,
	HEMBA1006865,	HEMBA1006877,	HEMBA1006885,	HEMBA1006900,	HEMBA1006914,
40	HEMBA1006921,	HEMBA1006926,	HEMBA1006936,	HEMBA1006941,	HEMBA1006949,
	HEMBA1006973,	HEMBA1006993,	HEMBA1007002,	HEMBA1007045,	HEMBA1007051,
	HEMBA1007052,	HEMBA1007062,	HEMBA1007066,	HEMBA1007073,	HEMBA1007078,
	HEMBA1007080,	HEMBA1007085, HEM	MBA1007087, HEMB	A1007113, HEMBA	1007121,
45	HEMBA1007129,	HEMBA1007147,	HEMBA1007149,	HEMBA1007151,	HEMBA1007174,
	HEMBA1007178,	HEMBA1007194,	HEMBA1007203,	HEMBA1007206,	HEMBA1007224,
	HEMBA1007243,	HEMBA1007256,	HEMBA1007267,	HEMBA1007273,	HEMBA1007279,
	HEMBA1007288,	HEMBA1007300,	HEMBA1007301,	HEMBA1007320,	HEMBA1007322,
<b>50</b>	HEMBA1007327,	HEMBA1007341,	HEMBA1007347,	HEMBB1000005,	HEMBB1000008,
50	HEMBB1000018,	HEMBB1000024,	HEMBB1000025,	HEMBB1000030,	HEMBB1000036,
	HEMBB1000037,	HEMBB1000039,	HEMBB1000044,	HEMBB1000050,	HEMBB1000054,
	HEMBB1000055,	HEMBB1000059,	HEMBB1000083,	HEMBB1000089,	HEMBB1000099,
	HEMBB1000103,	HEMBB1000113,	HEMBB1000119,	HEMBB1000136,	HEMBB1000141,
55	HEMBB1000144,	HEMBB1000173,	HEMBB1000198,	HEMBB1000215,	HEMBB1000217,

	HEMBB1000218,	HEMBB1000226,	HEMBB1000240,	HEMBB1000244.	HEMBB1000258,
	HEMBB1000264,	HEMBB1000266.	HEMBB1000274,	HEMBB1000307,	HEMBB1000312,
	HEMBB1000317,	HEMBB1000318,	HEMBB1000336.	HEMBB1000337.	HEMBB1000338,
5	HEMBB1000339,	HEMBB1000341,	HEMBB1000343,	HEMBB1000354,	HEMBB1000369,
	HEMBB1000374,	HEMBB1000376,	HEMBB1000391,	HEMBB1000402.	HEMBB1000420,
	HEMBB1000434,	HEMBB1000438,	HEMBB1000441,	HEMBB1000455,	HEMBB1000472.
	HEMBB1000480,	HEMBB1000490.	HEMBB1000491,	HEMBB1000493,	HEMBB1000510,
10	HEMBB1000523,	HEMBB1000530,	HEMBB1000550,	HEMBB1000554,	HEMBB1000556,
	HEMBB1000564.	HEMBB1000573,	HEMBB1000575,	HEMBB1000586,	HEMBB1000589,
	HEMBB1000591,	HEMBB1000593, HE	•	•	•
	HEMBB1000631,	HEMBB1000632,	HEMBB1000637,	HEMBB1000638,	HEMBB1000643,
15	HEMBB1000649,	HEMBB1000652,	HEMBB1000671,	HEMBB1000684,	HEMBB1000693,
	HEMBB1000705,	HEMBB1000706.	HEMBB1000709,	HEMBB1000726,	HEMBB1000738,
	HEMBB1000749,	HEMBB1000763,	HEMBB1000770,	HEMBB1000781,	HEMBB1000790,
	HEMBB1000807,	HEMBB1000810,	HEMBB1000821,	HEMBB1000826,	HEMBB1000827,
20	HEMBB1000835,	HEMBB1000840,	HEMBB1000848,	HEMBB1000870,	HEMBB1000887,
20	HEMBB1000890.	HEMBB1000893,	HEMBB1000908,	HEMBB1000910,	HEMBB1000913.
	HEMBB1000915,	HEMBB1000917,	HEMBB1000927,	HEMBB1000947,	HEMBB1000959.
	HEMBB1000975,	HEMBB1000991.	HEMBB1000996,	HEMBB1001014,	HEMBB1001020,
	HEMBB1001024,	HEMBB1001037,	HEMBB1001047,	HEMBB1001056,	HEMBB1001058;
25	HEMBB1001060,	HEMBB1001063,	HEMBB1001068,	HEMBB1001096,	HEMBB1001102,
	HEMBB1001105,	HEMBB1001112,	HEMBB1001114,	HEMBB1001119,	HEMBB1001126,
	HEMBB1001133,	HEMBB1001137,	HEMBB1001142,	HEMBB1001151,	HEMBB1001153,
	HEMBB1001169,	HEMBB1001175,	HEMBB1001177,	HEMBB1001182,	HEMBB1001208,
30	HEMBB1001209,	HEMBB1001210,	HEMBB1001218,	HEMBB1001234,	HEMBB1001249,
	HEMBB1001253,	HEMBB1001254,	HEMBB1001267,	HEMBB1001271,	HEMBB1001282,
	HEMBB1001288,	HEMBB1001289,	HEMBB1001294,	HEMBB1001302,	HEMBB1001317,
	HEMBB1001337,	HEMBB1001346,	HEMBB1001348,	HEMBB1001366,	HEMBB1001367,
35	HEMBB1001369,	HEMBB1001380,	HEMBB1001426,	HEMBB1001429,	HEMBB1001436,
	HEMBB1001449,	HEMBB1001454, HE	MBB1001458, HEMB	B1001463, HEMBB1	001500,
	HEMBB1001521,	HEMBB1001527,	HEMBB1001531,	HEMBB1001535,	HEMBB1001536,
	HEMBB1001537,	HEMBB1001555,	HEMBB1001562,	HEMBB1001564,	HEMBB1001565,
40	HEMBB1001585,	HEMBB1001586,	HEMBB1001588,	HEMBB1001618,	HEMBB1001619,
	HEMBB1001635,	HEMBB1001637,	HEMBB1001653,	HEMBB1001673,	HEMBB1001704,
	HEMBB1001706,	HEMBB1001707,	HEMBB1001717,	HEMBB1001735,	HEMBB1001749,
	HEMBB1001753,	HEMBB1001756,	HEMBB1001802,	HEMBB1001812,	HEMBB1001816,
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[0202] Genes that were expressed in all the tissues tested are indicated below by the corresponding clone names:

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Ū	PLACE1004358,	PLACE1004672,	PLACE1004736,	PLACE1004824,	PLACE1004900.
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	THYR01000186,	THYR01000484,	THYR01000569,	THYR01000699,	THYR01000712,
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	THYR01001411,	THYR01001426, TH	YR01001480, THYF	R01001487, THYR01	001537,
	THYR01001637,	THYR01001772,	THYR01001793,	THYR01001828,	THYR01001854,
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	Y79AA1000560,	Y79AA1000589,	Y79AA1000833,	Y79AA1000985,	Y79AA1001077,
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35	Y79AA1002482.				

[0203] Genes that were expressed at low levels in any of the tissues tested are indicated below by the corresponding clone names:

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	HEMBA1000302,	HEMBA1000307,	HEMBA1000333,	HEMBA1000390,	HEMBA1000418,
5	HEMBA1000434,	HEMBA1000442,	HEMBA1000460,	HEMBA1000464,	HEMBA1000490,
	HEMBA1000504,	HEMBA1000518,	HEMBA1000520,	HEMBA1000531,	HEMBA1000555,
	HEMBA1000563,	HEMBA1000592,	HEMBA1000594,	HEMBA1000608,	HEMBA1000637,
	HEMBA1000662,	HEMBA1000686,	HEMBA1000719,	HEMBA1000722,	HEMBA1000727,
10	HEMBA1000747,	HEMBA1000773,	HEMBA1000817,	HEMBA1000822,	HEMBA1000851,
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	HEMBA1001024,	HEMBA1001026,	HEMBA1001043,	HEMBA1001052,	HEMBA1001071,
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25	THYR01000129,	THYR01000156,	THYR01000199,	THYR01000242,	THYR01000270,
	THYR01000279,	THYR01000288,	THYR01000327,	THYR01000381,	THYR01000438;
	THYR01000488,	THYR01000501,	THYR01000502,	THYR01000505,	THYR01000570,
	THYR01000596,	THYR01000605, THY	/R01000641, THYR	01000662, THYR01	000734,
30	THYR01000777,	THYR01000783,	THYR01000793,	THYR01000796,	THYR01000805,
	THYR01000829,	THYR01000852,	THYR01000895,	THYR01000934,	THYR01000952,
	THYR01001033,	THYR01001100,	THYR01001134,	THYR01001142,	THYR01001204,
	THYR01001262,	THYR01001271,	THYR01001290,	THYR01001347,	THYR01001559,
35	THYRO1001661,	THYR01001721,	THYR01001745,	Y79AA1000328,	Y79AA1000420,
	Y79AA1000705,	Y79AA1000734,	Y79AA1000748,	Y79AA1000752,	Y79AA1000774,
	Y79AA1000784,	Y79AA1000802,	Y79AA1000824,	Y79AA1000827,	Y79AA1000976,
	Y79AA1001078,	Y79AA1001281,	Y79AA1001312,	Y79AA1001493,	Y79AA1001541,
40	Y79AA1001581,	Y79AA1001585,	Y79AA1001696,	Y79AA1001781,	Y79AA1001848,
	Y79AA1001874,	Y79AA1001923,	Y79AA1002027,	Y79AA1002115,	Y79AA1002139,
	Y79AA1002208,	Y79AA1002209,	Y79AA1002210,	Y79AA1002220,	Y79AA1002298,
	Y79AA1002307,	Y79AA1002311,	Y79AA1002407,	Y79AA1002416,	Y79AA1002431,
45	Y79AA1002487.				

[0204] Genes exhibiting characteristic features in the expression thereof were selected by statistical analysis of these data. Two examples are shown below to describe the selection of genes of which expression is varied greatly among tissues. The β' actin gene is used frequently as a control in gene expression analysis. Genes of which expression is varied greatly among tissues as compared that of the β-actin gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β-actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_a^2$ . Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_b^2$ . By taking variance ratio F as  $F=S_b^2/S_a^2$ , genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below by the corresponding clone names:

HEMBA1002113, HEMBA1005296, HEMBA1007121, HEMBB1000637, HEMBB1000915, MAMMA1000597, MAMMA1000605, MAMMA1000962, MAMMA1001139, MAMMA1001198, MAMMA1002858, NT2RM2001896, NT2RP2002710, NT2RP2004339, NT2RP2004538, NT2RP3000348, NT2RP3003121, PLACE3000009,

PLACE3000254, THYRO1000569, Y79AA1000131.

[0205] Gene of OVARC1000037 {heterogeneous nuclear ribonucleoprotein (hnRNP)} which expression is varied little. Genes of which expression is varied greatly among tissues as compared that of the OVARC1000037 gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of  $\beta$ -actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_a{}^2$ . Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance  $S_b{}^2$ . By taking variance ratio F as  $F = S_b{}^2/S_a{}^2$ , genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below,by the corresponding clone names:

40					
10	HEMBA1000304,	NT2RM2001716,	NT2RM2001840,	HEMBA1001051,	HEMBA1001109,
	OVARC1001731,	HEMBA1000726,	HEMBA1001286,	HEMBA1000387,	HEMBA1000519,
	NT2RM2001896,	HEMBA1000042,	HEMBA1001085,	HEMBA1001330,	OVARC1000576_
45	HEMBA1000575,	NT2RM2000599,	NT2RM2000714,	HEMBA1000469,	NT2RM4000366,
15	HEMBA1001377,	HEMBA1000769,	HEMBA1000338,	NT2RM2000795,	HEMBA1001299,
	HEMBA1000508,	HEMBA1000150,	HEMBA1000774,	HEMBA1001226,	HEMBA1000960;
	NT2RM4000795,	HEMBA1002162,	NT2RM4001876,	NT2RM4002482,	HEMBA1001678,
	HEMBA1002113,	NT2RM4002383,	HEMBA1002229,	HEMBA1002818,	HEMBA1001454,
20	NT2RM4000764,	HEMBA1001510,	HEMBA1001714,	HEMBA1002150,	NT2RM4002044,
	HEMBA1002728,	NT2RM4002189,	HEMBA1001991,	HEMBA1002166,	NT2RM4002499,
	NT2RM4001140,	NT2RM4002504,	HEMBA1002590,	HEMBA1001435,	PLACE1000706,
0.5	HEMBA1002160,	HEMBA1001824,	HEMBA1001463,	HEMBA1001533,	HEMBA1001570,
25	PLACE1001036,	HEMBA1001651,	HEMBA1002381,	HEMBA1002934,	HEMBA1003370,
	HEMBA1003021,	HEMBA1003166,	NT2RP1000738,	NT2RP2000040,	HEMBA1004164,
	HEMBA1003836,	HEMBA1004267,	NT2RP2000845,	HEMBA1003041,	HEMBA1003571,
20	HEMBA1003758,	NT2RP2000108,	HEMBA1003838,	NT2RP1000357,	HEMBA1003376,
30	PLACE1003528,	HEMBA1003528,	NT2RP1001475,	HEMBA1004049,	HEMBA1003212,
	HEMBA1003667,	PLACE1004149,	HEMBA1003926,	HEMBA1004306,	HEMBA1004024,
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25	NT2RP2001081,	HEMBA1004056, HE	MBA1003314, HEMB	A1003827, HEMBA1	003893,
35	NT2RP2001036,	NT2RP2001168,	NT2RP2001328,	HEMBA1005035,	NT2RP2001569,
	NT2RP2002439,	HEMBA1005511,	HEMBA1005999,	NT2RP2002862,	NT2RP2002979,
	NT2RP2001394,	HEMBA1004753,	NT2RP2002621,	HEMBA1005853,	HEMBA1005443,
40	NT2RP2002980,	NT2RP2001347,	HEMBA1005241,	NT2RP2002750,	NT2RP2003533,
40	HEMBA1005634,	NT2RP2003034,	HEMBA1006138,	NT2RP2003117,	NT2RP2001366,
	HEMBA1005079,	NT2RP2003293,	NT2RP2002710,	HEMBA1005911,	NT2RP2002752,
	HEMBA1006036,	NT2RP2002987,	HEMBA1006100,	HEMBA1004460,	HEMBA1004538,
45	NT2RP2001943,	NT2RP2002033,	HEMBA1005296,	HEMBA1005829,	HEMBA1005520,
45	HEMBA1005123,	HEMBA1005552,	HEMBA1004930,	NT2RP2001312,	HEMBA1005304,
	HEMBA1005834,	HEMBA1005990,	HEMBA1005526,	NT2RP2003073,	HEMBA1005331,
	HEMBA1006744,	HEMBA1006780,	NT2RP2004339,	HEMBB1000173,	HEMBA1007113,
50	NT2RP2005908,	HEMBB1000376,	HEMBB1000024,	HEMBB1000510,	NT2RP2004580,
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	NT2RP2003912,	HEMBA1006381,	NT2RP2004270,	HEMBA1006993,	NT2RP2005325,
	HEMBB1000218,	HEMBA1007121,	HEMBB1000434,	NT2RP2006166,	PLACE1010310.
	HEMBB1000589,	NT2RP2003986,	HEMBA1006419,	HEMBB1000141,	HEMBB1000030.
5	HEMBB1000099,	HEMBA1006624,	NT2RP2003988,	NT2RP2005753,	HEMBA1007147,
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	HEMBA1006631,	NT2RP2004538,	HEMBA1006695,	HEMBB1000490.	NT2RP2006184.
	HEMBB1000573,	NT2RP2004675,	HEMBA1007078,	NT2RP2005457,	NT2RP2005491.
10	HEMBB1000491,	HEMBB1000575, PLA	•	•	
	HEMBB1000337,	HEMBB1000008,	NT2RP2005701,	NT2RP2003825,	HEMBB1000215,
	HEMBB1000258,	NT2RP2005496,	HEMBB1000374,	NT2RP2005581,	HEMBB1000018.
	HEMBB1000493,	HEMBB1000554,	HEMBB1000671,	NT2RP3000348,	HEMBB1001267,
15	HEMBB1002092,	HEMBB1001749,	PLACE3000304,	HEMBB1001834,	HEMBB1001142,
	HEMBB1001177,	HEMBB1000840,	HEMBB1001436,	HEMBB1001921,	HEMBB1002005,
	NT2RP3001398,	HEMBB1002069,	HEMBB1002094,	HEMBB1001836,	HEMBB1001114,
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	NT2RP2006571,	HEMBB1001802,	HEMBB1000915,	HEMBB1000996,	NT2RP3001274,
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	HEMBB1000917,	PLACE1011090,	NT2RP3000742,	PLACE2000435,	HEMBB1001944,
	PLACE3000254,	HEMBB1002218,	NT2RP2006436,	NT2RP3000320,	HEMBB1001126,
	HEMBB1000749,	NT2RP3000644,	HEMBB1001527,	NT2RP3000850,	HEMBB1001967;
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	MAMMA1000348,	MAMMA1000372,	NT2RP3003121,	MAMMA1000851,	MAMMA1000501,
30	THYRO1000569,	MAMMA1000576,	MAMMA1000605,	HEMBB1002442,	HEMBB1002596,
	MAMMA1000198,	MAMMA1000714,	MAMMA1000356,	MAMMA1000760,	MAMMA1000416,
	MAMMA1000931,	MAMMA1000962,	HEMBB1002387,	NT2RP3001712,	HEMBB1002556,
	HEMBB1002617,	MAMMA1000171, PLA	CE4000612, MAM	MA1000761, MAMMAI	.000421,
35	MAMMA1000444,	HEMBB1002358,	HEMBB1002453,	HEMBB1002664,	HEMBB1002699,
	MAMMA1000720,	MAMMA1000360,	MAMMA1000744,	MAMMA1000802,	MAMMA1000839,
	MAMMA1000478,	MAMMA1000877,	MAMMA1000585,	MAMMA1000941,	NT2RP3002909,
	MAMMA1000968,	HEMBB1002457,	HEMBB1002495,	MAMMA1000057,	MAMMA1000696,
40	MAMMA1000942,	MAMMA1000302,	MAMMA1000943,	MAMMA1000998,	NT2RP3001752,
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	NT2RP3002146,	MAMMA1000084,	MAMMA1000257,	NT2RP3003157,	MAMMA1000431,
	MAMMA1000597,	MAMMA1001078,	MAMMA1001126,	THYR01001537,	NT2RP3004093,
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45	Y79AA1000560,	MAMMA1002359,	MAMMA1002056,	MAMMA1002413,	MAMMA1002454,
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	MAMMA1001745,	MAMMA1001790,	NT2RP3004470,	NT2RP3004503,	MAMMA1002293,
50	NT2RP4000321,	MAMMA1001880,	Y79AA1000410,	MAMMA1001970,	MAMMA1002058,
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	MAMMA1002032,	MAMMA1002125,	MAMMA1002174,	NT2RP4000614,	NT2RP4000243,
55	NT2RP3004110,	MAMMA1001271,	MAMMA1001296,	MAMMA1001992,	MAMMA1002033,

	MAMMA1002428,	MAMMA1002590,	MAMMA1001186,	MAMMA1002267,	MAMMA1002322,
	MAMMA1001956,	MAMMA1002155, NT	2RP4000210, MAMMA	1002622, NT2RP30	004125,
5	MAMMA1001220,	MAMMA1001683,	NT2RP3004348,	Y79AA1000214,	Y79AA1000833,
5	NT2RP4000212,	MAMMA1002230,	MAMMA1001452,	MAMMA1001620,	MAMMA1001256,
	MAMMA1001760,	NT2RP3004349,	MAMMA1001783,	MAMMA1001907,	MAMMA1002009,
	MAMMA1002545,	NT2RP4000214,	NT2RP4000728,	MAMMA1001465,	MAMMA1001154,
10	MAMMA1001198,	MAMMA1001343,	MAMMA1002310,	NT2RP4000035,	NT2RP4000833,
10	MAMMA1003150,	MAMMA1002886,	NT2RP4001938,	NT2RM2000260,	MAMMA1002629,
	MAMMA1002973,	MAMMA1002721,	MAMMA1002909,	NT2RP4001100,	NT2RM1000857,
	NT2RP4000878,	MAMMA1002844,	NT2RM1000039,	NT2RP4001174,	MAMMA1002665,
	MAMMA1003047,	NT2RM1000086,	NT2RM1000260,	NT2RM1000355,	MAMMA1002701,
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	MAMMA1003004,	MAMMA1002673,	MAMMA1003031,	MAMMA1002764,	MAMMA1002858,
	NT2RP4001679,	NT2RP4002888,	MAMMA1002711,	NT2RP4001276,	NT2RM1000018,
	NT2RP4001568,	NT2RM1000883.			

[0206] Thus, characteristic features in the expression of a gene are illustrated by comparing and \_ statistically analyzing the expression of many genes.

### 5 Analysis of disease-associated genes

[0207] Non-enzymic protein glycation reaction is believed to be a cause of a variety of chronic diabetic complications. Accordingly, genes of which expression is elevated or decreased in a glycated protein-specific manner in the endothelial cells are associated with diabetic complications caused by glycated proteins. Vascular endothelial cells are affected with glycated proteins present in blood. Reaction products of non-enzymic protein glycation include amadori compound (glycated protein) as a mildly glycated protein and advanced glycation endproduct as a heavily glycated protein. Hence, a survey was carried out for genes of which expression levels are varied depending on the presence of these glycated proteins in endothelial cells. The mRNAs were extracted from endothelial cells that were cultured in the presence or absence of glycated protein. The mRNAs were converted into radiolabeled first strand cDNAs for preparing probes. The probes were hybridized to the above-mentioned DNA array. Signal of each DNA spot was detected by BAS2000 and analyzed by ArrayGauge (Fuji Photo Film Co., Ltd.).

[0208] Advanced glycation endproduct of bovine serum albumin was prepared as follows: bovine serum albumin (BSA; Sigma) was incubated in a phosphate buffer solution containing 50 mM glucose at 37°C for 8 weeks; and the resulting brownish BSA was dialyzed against a phosphate buffer solution.

[0209] Human normal pulmonary arterial endothelial cells (Cell Applications) were cultured in an Endothelial Cell Growth Medium (Cell Applications). The culture dish (Farcon) with the cells were incubated in a CO<sub>2</sub> incubator (37°C, 5% CO<sub>2</sub>, in a humid atmosphere). When the cells were grown to be confluent in the dish, 250 μg/ml of bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin was added thereto and the cells were incubated for 33 hours. The mRNA was extracted from the cells by using a FastTack<sup>(TM)</sup> 2.0 kit (Invitrogen). The labeling of hybridization probe was carried out by using the mRNA according to the same procedure as described above.

[0210] Table 169 shows the expression level of each cDNA in human pulmonary arterial endothelial cells cultured in a medium containing bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin. Genes of which expression was detected in the endothelial cell are as follows:

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	HEMBA1000005,	HEMBA1000012,	HEMBA1000020,	HEMBA1000042,	HEMBA1000046,
	HEMBA1000076,	HEMBA1000111,	HEMBA1000150,	HEMBA1000185,	HEMBA1000216.
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5	HEMBA1000327,	HEMBA1000338,	HEMBA1000356,	HEMBA1000357.	HEMBA1000366.
	HEMBA1000376,	HEMBA1000387,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000460,	HEMBA1000469.	HEMBA1000501,	HEMBA1000504,	HEMBA1000505,
	HEMBA1000508,	HEMBA1000519,	HEMBA1000534,	HEMBA1000540,	
10	HEMBA1000555,	HEMBA1000557.	HEMBA1000575,	HEMBA1000592,	HEMBA1000542,
70	HEMBA1000622,	HEMBA1000636,	HEMBA1000575,	HEMBA1000532,	HEMBA1000604,
	HEMBA1000686.	HEMBA1000702,			HEMBA1000682,
	HEMBA1000769,	HEMBA1000774.	HEMBA1000726,	HEMBA1000727,	HEMBA1000749,
	HEMBA1000872,		HEMBA1000791,	HEMBA1000827,	HEMBA1000852,
15	HEMBA1000974,	HEMBA1000876,	HEMBA1000918,	HEMBA1000942,	HEMBA1000960,
	HEMBA1001080,	HEMBA1001020,	HEMBA1001051,	HEMBA1001059,	HEMBA1001060,
		HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001122,
	HEMBA1001123,	HEMBA1001140,	HEMBA1001172,	HEMBA1001196,	HEMBA1001197,
20	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,	HEMBA1001247,	HEMBA1001265,
	HEMBA1001281,	HEMBA1001286,	HEMBA1001299,	HEMBA1001302,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,	HEMBA1001377,	HEMBA1001387,	HEMBA1001398,
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	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,	HEMBA1001595,	HEMBA1001608,
	HEMBA1001620,	HEMBA1001640,	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,
	HEMBA1001658,	HEMBA1001678,	HEMBA1001714,	HEMBA1001718,	HEMBA1001723,
30	HEMBA1001746,	HEMBA1001784,	HEMBA1001791,	HEMBA1001800,	HEMBA1001804,
	HEMBA1001809,	HEMBA1001819,	HEMBA1001824,	HEMBA1001844,	HEMBA1001866,
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	HEMBA1001960,	HEMBA1001987,	HEMBA1001991,	HEMBA1002003,	HEMBA1002008,
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	HEMBA1002337,	HEMBA1002348,	HEMBA1002381,	HEMBA1002458,	HEMBA1002462,
40	HEMBA1002469,	HEMBA1002503,	HEMBA1002508,	HEMBA1002542,	HEMBA1002547,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002583,	HEMBA1002590,	HEMBA1002592,
	HEMBA1002609,	HEMBA1002624,	HEMBA1002629,	HEMBA1002645,	HEMBA1002678,
	HEMBA1002688,	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,
45	HEMBA1002750,	HEMBA1002770,	HEMBA1002779,	HEMBA1002780,	HEMBA1002818,
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	HEMBA1003142,	HEMBA1003166,	HEMBA1003175,	HEMBA1003179,	HEMBA1003202,
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	HEMBA1003640,	HEMBA1003656,	HEMBA1003667,	HEMBA1003680,	HEMBA1003690,
5	HEMBA1003692,	HEMBA1003715,	HEMBA1003720,	HEMBA1003742,	HEMBA1003758,
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55	MAMMA1000302,	MAMMA1000307,	MAMMA1000279,	MAMMA1000254,	MAMMA1000360,
		MARGINIA TOOLOGO (	יייוווותנייויי,	INTERNITY TO CO 200'	יייטסרטטר דערמיייייייייייייייייייייייייייייייייייי

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[0211] Signal ratios of EC\_AGE\_BSA to EC\_BSA and of EC\_glycated\_BSA to EC\_BSA were calculated for each gene. Genes with high signal ratios were selected. In the case of calculating the ratio of signal value of 40 or less to that of more than 40, such signal values were, for convenience, taken as 40 instead of the real values. When the ratio EC\_AGE\_BSA/EC\_BSA is 2 or more, expression of the genes exhibiting such ratio is expected to be elevated due to advanced glycation endproduct of bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC\_AGE\_BSA/EC\_BSA ranges from 0.5 to 2, expression of the genes exhibiting such ratio is expected to be unaffected due to advanced glycation endproduct of bovine serum albumin. When the ratio EC\_AGE\_BSA/EC\_BSA is less than 0.5, expression of the genes exhibiting such ratio value is expected to be decreased due to advanced glycation endproduct of bovine serum albumin. The lower the value is, the lower the gene expression level is. [0212] Clone with EC\_AGE\_BSA/EC\_BSA ratio of 2 or higher are as follows: HEMBA1003958, MAMMA1001256, PLACE2000411.

[0213] Clone with EC\_AGE\_BSA/EC\_BSA ratio of 0.5 or less is as follows: MAMMA1001783. These cDNAs are associated with diabetes.

[0214] When the ratio EC\_glycated\_BSA/EC\_BSA is 2 or more, the expression level of the gene exhibiting such ratio is expected to be elevated due to glycated bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC\_glycated\_BSA/EC\_BSA ranges from 0.5 to 2, the expression level of the gene exhibiting such ratio is expected to be unaffected with glycated bovine serum albumin. When the ratio EC\_glycated\_BSA/EC\_BSA is less than 0.5, the expression level of a gene exhibiting such ratio is expected to be decreased due to glycated bovine serum albumin. The lower the value is, the lower the gene expression level is.

[0215] Clones with EC\_glycated\_BSA/EC\_BSA ratio of 2 or more are as follows: HEMBA1004850, MAMMA1001256, MAMMA1002132 and PLACE3000119.

[0216] A clone with EC\_glycated\_BSA/EC\_BSA ratio of 0.5 or less is as follows: MAMMA1001783.

[0217] These cDNAs are also associated with diabetes.

4

Analysis of genes associated with neural cell differentiation

[0218] Genes involved in neural cell differentiation are useful for treating neurological diseases. It is possible that genes with varying expression levels in response to induction of cellular differentiation in neural cells are associated with neurological diseases.

[0219] A survey was performed for genes of which expression levels are varied in response to induction of differentiation (stimulation by retinoic acid (RA)) in cultured cells of a neural strain, NT2.

**[0220]** The NT2 cells were treated basically according to supplier's instruction manual. "Undifferentiated NT2 cells" means NT2 cells successively cultured in an Opti-MEM I (GIBCO-BRL; catalog No. 31985) containing 10%(v/v) fetal bovine serum and 1%(v/v) penicillin-streptomycin (GIBCO BRL). "NT2 cells cultured in the presence of retinoic acid" means the cells resulted from transferring undifferentiated NT2 cells into a retinoic acid-containing medium, which consists of D-MEM (GIBCO BRL; catalog No. 11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin and  $10~\mu$ M retinoic acid (GIBCO-BRL), and the subsequent successive culture therein for 5 weeks. "NT2 cells that were cultured in the presence of retinoic acid and then further cultured in the presence of cell-division inhibitor added" means NT2 cells resulted from transferring NT2 cells cultured in the presence of retinoic acid for 5 weeks into a cell-division inhibitor-containing medium, which consisted of D-MEM(GIBCO BRL; catalog No.11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin,  $10\mu$ M retinoic acid,  $10~\mu$ M FudR (5-fluoro-2'-deoxyuridine: GIBCO BRL),  $10~\mu$ M Urd (Uridine: GIBCO BRL) and  $1~\mu$ M araC (Cytosine β-D-Arabinofuranoside: GIBCO BRL), and the subsequence succes-

sive culture for 2 weeks. Each of the cells were treated with trypsin and then harvested. Total RNAs were extracted from the cells by using S.N.A.P.<sup>(TM)</sup> Total RNA Isolation kit (Invitrogen<sup>(r)</sup>). The labeling of probe used for hybridization was carried out by using 10 μg of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of stimulation for inducing differentiation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment-of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis, clones with the difference can be statistically detected even when the signals were low. Accordingly, clones with signal value of 40 or less were also assessed for the selection. [0221] Tables 170-349 show the expression level of each cDNA in undifferentiated NT2 cells, NT2 cells cultured in the presence of RA, and NT2 cells that were cultured in the presence of cell-division inhibitor added.

[0222] Averaged signal values  $(M_1, M_2)$  and sample variances  $(s_1^2, s_2^2)$  were calculated for each gene in each of the cells, and then, the pooled sample variances  $s^2$  were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula:  $t=(M1-M2)/s/(1/3+1/3)^{1/2}$ . When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information on an increase (+) or decrease (-) in the expression level of a gene in the treated cells when the level is compared with that of untreated undifferentiated cells.

[0223] Clones of which expression levels increased by RA are as follows: HEMBA1000005, HEMBA1000042, HEMBA1000046. HEMBA1000076. HEMBA1000111, HEMBA1000141, HEMBA1000150, HEMBA1000185. HEMBA1000282, HEMBA1000304, HEMBA1000307, HEMBA1000338, HEMBA1000357, HEMBA1000376, HEMBA1000387. HEMBA1000392, HEMBA1000428, HEMBA1000456, HEMBA1000459, HEMBA1000469. HEMBA1000504, HEMBA1000508, HEMBA1000519, HEMBA1000540, HEMBA1000545. HEMBA1000557, HEMBA1000563, HEMBA1000568, HEMBA1000575. HEMBA1000588, HEMBA1000592, HEMBA1000604, HEMBA1000622, HEMBA1000655, HEMBA1000673, HEMBA1000682, HEMBA1000726. HEMBA1000727, HEMBA1000749, HEMBA1000769, HEMBA1000774, HEMBA1000791, HEMBA1000822, HEMBA1000872, HEMBA1000876. HEMBA1000910, HEMBA1000942, HEMBA1000943, HEMBA1000960, HEMBA1000972, HEMBA1000974, HEMBA1000991, HEMBA1001008, HEMBA1001020, HEMBA1001043, HEMBA1001051, HEMBA1001060, HEMBA1001071, HEMBA1001077, HEMBA1001085, HEMBA1001094, HEMBA1001109, HEMBA1001121, HEMBA1001122, HEMBA1001140. HEMBA1001172. HEMBA1001226, HEMBA1001235, HEMBA1001265, HEMBA1001281, HEMBA1001294, HEMBA1001299, HEMBA1001319. HEMBA1001323, HEMBA1001351, HEMBA1001330, HEMBA1001361, HEMBA1001377, HEMBA1001388, HEMBA1001391. HEMBA1001398, HEMBA1001432, HEMBA1001435, HEMBA1001442, HEMBA1001497, HEMBA1001454, HEMBA1001455. HEMBA1001517, HEMBA1001569, HEMBA1001570, HEMBA1001581. HEMBA1001585, HEMBA1001620, HEMBA1001711, HEMBA1001718, HEMBA1001723, HEMBA1001761. HEMBA1001815. HEMBA1001819, HEMBA1001864, HEMBA1001861, HEMBA1001869, HEMBA1001888. HEMBA1001915, HEMBA1001918, HEMBA1001940, HEMBA1001964, HEMBA1001967, HEMBA1001979, HEMBA1001987, HEMBA1001991, HEMBA1002008, HEMBA1002022, HEMBA1002039, HEMBA1002049. HEMBA1002084, HEMBA1002102. HEMBA1002113, HEMBA1002144, HEMBA1002160. HEMBA1002162. HEMBA1002185. HEMBA1002212. HEMBA1002226, HEMBA1002229, HEMBA1002267. HEMBA1002270, HEMBA1002337, HEMBA1002381. HEMBA1002458, HEMBA1002477, HEMBA1002508, HEMBA1002558, HEMBA1002561, HEMBA1002583. HEMBA1002590, HEMBA1002628, HEMBA1002645, HEMBA1002661, HEMBA1002678, HEMBA1002712, HEMBA1002728. HEMBA1002780. HEMBA1002850. 45 HEMBA1002886, HEMBA1002934, HEMBA1002935. HEMBA1002939. HEMBA1002951. HEMBA1002968. HEMBA1002970, HEMBA1002973, HEMBA1002999, HEMBA1003021, HEMBA1003033, HEMBA1003034, HEMBA1003064, HEMBA1003067, HEMBA1003078. HEMBA1003086, HEMBA1003129, HEMBA1003096, HEMBA1003142, HEMBA1003148, HEMBA1003166, HEMBA1003175, HEMBA1003197, HEMBA1003199, HEMBA1003202, HEMBA1003204, HEMBA1003212, HEMBA1003235, HEMBA1003250, HEMBA1003273, HEMBA1003276, HEMBA1003278. HEMBA1003291, HEMBA1003309, HEMBA1003322, HEMBA1003328, HEMBA1003348, HEMBA1003369, HEMBA1003376, HEMBA1003463, HEMBA1003384, HEMBA1003395, HEMBA1003480. HEMBA1003531, HEMBA1003548, HEMBA1003591, HEMBA1003595, HEMBA1003597, HEMBA1003617, HEMBA1003621, HEMBA1003622, HEMBA1003637, HEMBA1003640, HEMBA1003645, HEMBA1003646, HEMBA1003656, HEMBA1003679, HEMBA1003692, HEMBA1003715, HEMBA1003720, HEMBA1003725, HEMBA1003729, HEMBA1003758, HEMBA1003803, HEMBA1003805, HEMBA1003836, HEMBA1003838, HEMBA1003879, HEMBA1003885, HEMBA1003893, HEMBA1003908, HEMBA1003937, HEMBA1003942, HEMBA1003953, HEMBA1003958, HEMBA1003959. HEMBA1003978, HEMBA1003987, HEMBA1003989, HEMBA1004000, HEMBA1004011. HEMBA1004012, HEMBA1004015, HEMBA1004024,

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    [0224] Clones of which expression levels decreased by RA are as follows:
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    [0225] Clones of which expression levels increase by RA/inhibitor are as follows:
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    Y79AA1002258.
    [0226] Clones of which expression levels decrease by RA/inhibitor are as follows:
    HEMBA1000012, HEMBA1000501, HEMBA1000946, HEMBA1003220, HEMBA1003403, HEMBA1003569,
    HEMBA1003591.
                   HEMBA1003926,
                                                                                 HEMBA1005296,
                                   HEMBA1004168,
                                                  HEMBA1004507,
                                                                 HEMBA1005009,
    HEMBA1005528, HEMBA1005570,
                                   HEMBA1006467,
                                                  HEMBA1006486,
                                                                 HEMBA1006492,
                                                                                 HEMBA1007322,
    HEMBB1000055, HEMBB1000244,
                                  HEMBB1001665, MAMMA1000684,
                                                                 MAMMA1001139,
                                                                                MAMMA1001743,
    NT2RM1000257,
                   NT2RM1000318,
                                   NT2RM1000539,
                                                  NT2RM1000666,
                                                                  NT2RM2000092,
                                                                                 NT2RM2000192,
    NT2RM2000371,
                   NT2RM2000594,
                                   NT2RM4000511,
                                                  NT2RM4001140,
                                                                  NT2RM4001754,
                                                                                 NT2RM4001905,
    NT2RM4001940,
                    NT2RM4002593,
                                   NT2RP1000086,
                                                  NT2RP1000439,
                                                                  NT2RP1001073,
                                                                                 NT2RP2000098,
    NT2RP2000965,
                   NT2RP2001397,
                                   NT2RP2002047,
                                                  NT2RP2004226,
                                                                  NT2RP2004396,
                                                                                 NT2RP2004655,
    NT2RP2005126,
                   NT2RP2005464,
                                   NT2RP2005712,
                                                  NT2RP2005859,
                                                                  NT2RP2005890,
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    NT2RP3001383,
                   NT2RP3001621,
                                   NT2RP3002081,
                                                  NT2RP3002181,
                                                                  NT2RP3002244,
                                                                                 NT2RP3002590,
    NT2RP3003059.
                   NT2RP3004258,
                                   NT2RP3004378.
                                                  NT2RP3004527.
                                                                  NT2RP3004594.
                                                                                 NT2RP4001760,
    NT2RP4001950,
                   NT2RP4002047.
                                   NT2RP4002408.
                                                  NT2RP5003459,
                                                                 OVARC1000004.
                                                                                 OVARC1000035.
35
    OVARC1000431,
                   OVARC1001051.
                                   OVARC1001129,
                                                  OVARC1001176,
                                                                  OVARC1001261,
                                                                                 OVARC1001342.
    OVARC1001942.
                    OVARC1001943.
                                   PLACE1002171.
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                                   PLACE1005876,
    PLACE1004128.
                   PLACE1005026.
                                                  PLACE1005923,
                                                                  PLACE1007257,
                                                                                 PLACE1007375.
    PLACE1007507.
                   PLACE1008941.
                                   PLACE1010624,
                                                  PLACE1011090,
                                                                  PLACE1011219,
                                                                                 THYRO1000270.
    Y79AA1000346, Y79AA1001541.
    [0227] Clones of which expression levels increase in the presence of both RA and RA/inhibitor are as follows:
    HEMBA1000046, HEMBA1000307, HEMBA1000504, HEMBA1000588, HEMBA1000682, HEMBA1000726,
    HEMBA1000943, HEMBA1001071, HEMBA1001094, HEMBA1001122,
                                                                 HEMBA1001323, HEMBA1001361,
    HEMBA1001455, HEMBA1001869,
                                  HEMBA1002084, HEMBA1002583.
                                                                 HEMBA1002628.
                                                                                 HEMBA1003096,
    HEMBA1003142, HEMBA1003276,
                                  HEMBA1003309, HEMBA1003463,
                                                                 HEMBA1003597,
                                                                                 HEMBA1003617,
45
    HEMBA1003725, HEMBA1003803,
                                  HEMBA1003879,
                                                  HEMBA1003989.
                                                                 HEMBA1004000,
                                                                                 HEMBA1004015,
    HEMBA1004024, HEMBA1004049,
                                  HEMBA1004056,
                                                                 HEMBA1004248,
                                                                                 HEMBA1004356,
                                                  HEMBA1004199,
    HEMBA1004666, HEMBA1004770,
                                  HEMBA1004803,
                                                  HEMBA1004923,
                                                                 HEMBA1004934,
                                                                                 HEMBA1004954,
    HEMBA1005039, HEMBA1005075,
                                                                                 HEMBA1005251,
                                   HEMBA1005113,
                                                  HEMBA1005219,
                                                                 HEMBA1005232,
    HEMBA1005304, HEMBA1005367,
                                   HEMBA1005410.
                                                  HEMBA1005411,
                                                                 HEMBA1005548.
                                                                                 HEMBA1005581,
    HEMBA1005631, HEMBA1005666.
                                  HEMBA1005780,
                                                  HEMBA1006485,
                                                                 HEMBA1006559,
                                                                                 HEMBA1006579.
    HEMBA1006754, HEMBB1000059.
                                                                                 HEMBB1000985.
                                  HEMBB1000575,
                                                  HEMBB1000709,
                                                                 HEMBB1000848,
    HEMBB1001117, HEMBB1001210,
                                  HEMBB1001394,
                                                                                 HEMBB1002049,
                                                  HEMBB1001668,
                                                                 HEMBB1001695,
    HEMBB1002254, HEMBB1002266,
                                  HEMBB1002371,
                                                                                 HEMBB1002697,
                                                 HEMBB1002614,
                                                                 HEMBB1002617,
    MAMMA1000241, MAMMA1000424, MAMMA1000616, MAMMA1000731, MAMMA1000824, MAMMA1001038,
    MAMMA1001243, MAMMA1002769, MAMMA1002871, MAMMA1002941,
    NT2RM1000725, NT2RM1000829,
                                  NT2RM2000504, NT2RM2000635,
                                                                  NT2RM2000821,
                                                                                 NT2RM2001582,
    NT2RM2001592,
                   NT2RM2001632,
                                   NT2RM2001648,
                                                  NT2RM2001671,
                                                                  NT2RM2001695,
                                                                                 NT2RM2001879,
    NT2RM4000290, NT2RM4000425,
                                  NT2RM4000471, NT2RM4000531,
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NT2RM4001454,
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                                                NT2RP1001013.
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NT2RP2002046,
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NT2RP2002701,
                NT2RP2003099,
                                NT2RP2003137,
                                                NT2RP2003206,
                                                                NT2RP2003230,
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NT2RP2003280,
                NT2RP2003393,
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                                                NT2RP2003456,
                                                                NT2RP2003596,
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NT2RP2004681,
                NT2RP2004743,
                                NT2RP2004775,
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                                                                NT2RP2004967,
                                                                                NT2RP2005003,
NT2RP2005289,
                NT2RP2005453.
                                NT2RP2005555,
                                                NT2RP2005767,
                                                                NT2RP2005853,
                                                                                NT2RP2006043,
NT2RP2006393,
                NT2RP2006436.
                                                                NT2RP2006565,
                                NT2RP2006441,
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NT2RP3000366,
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NT2RP3001007,
                NT2RP3001126,
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                                NT2RP3001782,
NT2RP3001679,
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NT2RP3002248,
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                                                                NT2RP3002549,
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NT2RP3003032,
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                                NT2RP3003193,
                                                NT2RP3003204,
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                NT2RP3003282.
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                                                NT2RP3004041,
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                NT2RP4000360,
NT2RP4000259.
                                NT2RP4000448,
                                                NT2RP4000524,
                                                                NT2RP4001079,
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NT2RP4001274,
                               NT2RP4001547,
                                               NT2RP4001677,
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OVARC1000384,
                OVARC1000408.
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                                               OVARC1000520,
                                                               OVARC1000526,
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OVARC1000679.
               OVARC1000682,
                                OVARC1000769,
                                               OVARC1000862,
                                                               OVARC1000984,
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OVARC1001004,
                OVARC1001154,
                                OVARC1001170,
                                               OVARC1001173,
                                                               OVARC1001200,
                                                                               OVARC1001268,
OVARC1001376, OVARC1001476, OVARC1001542, OVARC1001873, OVARC1001987, OVARC1002082,
PLACE1000014,
                PLACE1000048,
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                                               PLACE1000246,
                                                                PLACE1000292,
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PLACE1000347,
                PLACE1000712,
                                PLACE1001000,
                                               PLACE1001294,
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PLACE1001570,
                PLACE1001608,
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PLACE1001897,
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                                               PLACE1002157,
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                                                                                PLACE1002256,
PLACE1002259,
                PLACE1002438,
                               PLACE1002474,
                                               PLACE1002477,
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PLACE1002815,
                PLACE1002851,
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PLACE1003256,
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PLACE1003771,
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PLACE1004629,
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PLACE1006795.
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PLACE1007544,
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PLACE2000347,
               PLACE2000379,
                               PLACE3000121,
                                               PLACE3000124.
                                                                PLACE3000242.
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PLACE3000362,
               PLACE3000365,
                               PLACE3000400.
                                               PLACE3000401.
                                                               PLACE4000034.
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PLACE4000522,
               PLACE4000558,
                               THYRO1000197,
                                               THYRO1000241,
                                                               THYRO1000327,
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THYRO1000488,
               THYRO1000585,
                               THYRO1000625,
                                               THYRO1000805.
                                                               THYRO1000934,
                                                                               THYRO1001133,
THYRO1001173,
               THYRO1001213,
                               THYRO1001262,
                                                THYRO1001290,
                                                                Y79AA1000800,
                                                                                Y79AA1000976,
Y79AA1001585, Y79AA1001696, Y79AA1001711, Y79AA1002258.
[0228] Clones of which expression levels decrease in the presence of both RA and RA/inhibitor are as follows:
HEMBA1000946, HEMBA1003569, HEMBA1005570, NT2RM1000666,
                                                               NT2RM2000092,
                                                                               NT2RM2000594,
NT2RM4001754.
               NT2RM4001905,
                               NT2RP2002047, NT2RP3000980,
                                                               NT2RP3002081,
                                                                               NT2RP3004594,
NT2RP4001950.
               NT2RP4002408,
                               OVARC1000431, OVARC1001942,
                                                               OVARC1001943,
                                                                               PLACE1003190,
PLACE1005923, PLACE1007257, PLACE1010624, Y79AA1000346.
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55 [0229] These are neurological disease-associated clones.

#### Analysis of rheumatoid arthritis-associated genes

[0230] The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)-α participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis.

[0231] A survey was performed for genes of which expression levels are varied in response to TNF- $\alpha$  in the primary cell culture of synovial tissue. The primary cultured cells of the smooth muscle (Cell Applications) were grown to be confluent in a culture dish, and then, human TNF- $\alpha$  (Boehringer-Mannheim) was added at a final concentration of 10 ng/ml thereto. The culture was further continued for 24 hours.

[0232] Total RNA was extracted from the cells by using S.N.A.P.(TM) Total RNA Isolation kit (Invitrogen). The labeling of probe used for hybridization was carried out by using 10  $\mu$ g of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of TNF stimulation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis,

clones with the difference can be statistically detected even when the signals were low.

Accordingly, clones with signal value of 40 or less were also assessed for the selection.

Table 352 shows the expression level of each cDNA in synovial cells cultured in the absence or presence of TNF. [0233] Averaged signal values  $(M_1, M_2)$  and sample variances  $(s_1^2, s_2^2)$  for each gene were calculated in each of the cells, and then, the pooled sample variances  $s^2$  were obtained from the sample variances of the two types of cells to be compared. The t-values were determined according to the following formula:  $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}$ . When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively.

The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the stimulated cells when the level is compared with that of unstimulated cells.

[0234] Clones of which expression levels are elevated by TNF- $\alpha$  are as follows: HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000046, HEMBA1000076, HEMBA1000111, HEMBA1000168, HEMBA1000185. HEMBA1000201, HEMBA1000231. HEMBA1000243. HEMBA1000280. HEMBA1000282. HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000356, HEMBA1000376, HEMBA1000390, HEMBA1000387, HEMBA1000418, HEMBA1000460, HEMBA1000491, HEMBA1000501, HEMBA1000518, HEMBA1000519. HEMBA1000520, HEMBA1000531, HEMBA1000534, HEMBA1000542. HEMBA1000592, HEMBA1000545. HEMBA1000591. HEMBA1000594, HEMBA1000636, HEMBA1000655, HEMBA1000657. HEMBA1000673. HEMBA1000682. HEMBA1000686, HEMBA1000722, HEMBA1000726, HEMBA1000827, HEMBA1000870, HEMBA1000918, HEMBA1000971, HEMBA1000974, HEMBA1000986, HEMBA1001019, HEMBA1001043. HEMBA1001051, HEMBA1001059, HEMBA1001060, HEMBA1001071, HEMBA1001080. HEMBA1001109, HEMBA1001140, HEMBA1001172, HEMBA1001213, HEMBA1001196, HEMBA1001226, HEMBA1001281. HEMBA1001299, HEMBA1001302. HEMBA1001303, HEMBA1001323. HEMBA1001326. HEMBA1001327. HEMBA1001330. HEMBA1001351, HEMBA1001407, HEMBA1001411. HEMBA1001446, HEMBA1001454, HEMBA1001569. HEMBA1001647, HEMBA1001714, HEMBA1001800, HEMBA1001804, HEMBA1001809, HEMBA1001888, HEMBA1001912. HEMBA1001921, HEMBA1001967. 45 HEMBA1002084, HEMBA1002161, HEMBA1002166. HEMBA1002241. HEMBA1002337. HEMBA1002363. HEMBA1002389, HEMBA1002458, HEMBA1002460, HEMBA1002469, HEMBA1002538, HEMBA1002542, HEMBA1002547, HEMBA1002609, HEMBA1002624, HEMBA1002659, HEMBA1002750, HEMBA1002770, HEMBA1002779, HEMBA1002810, HEMBA1002816, HEMBA1002818, HEMBA1002850, HEMBA1002863, HEMBA1003021, HEMBA1003033, HEMBA1003078, HEMBA1003166, HEMBA1003202, HEMBA1003204, HEMBA1003229, HEMBA1003235, HEMBA1003276, HEMBA1003286, HEMBA1003296, HEMBA1003370, HEMBA1003376, HEMBA1003403. HEMBA1003418, HEMBA1003433, HEMBA1003447, HEMBA1003560, HEMBA1003569. HEMBA1003571, HEMBA1003591, HEMBA1003597, HEMBA1003598, HEMBA1003621, HEMBA1003656, HEMBA1003662, HEMBA1003680, HEMBA1003715, HEMBA1003725, HEMBA1003729, HEMBA1003733, HEMBA1003742, HEMBA1003773, HEMBA1003783, HEMBA1003950, HEMBA1004012, HEMBA1004015, HEMBA1004048, HEMBA1004074, HEMBA1004086, HEMBA1004111, HEMBA1004131. HEMBA1004202, HEMBA1004203, HEMBA1004207, HEMBA1004248, HEMBA1004274, HEMBA1004321, HEMBA1004330, HEMBA1004356, HEMBA1004366, HEMBA1004405, HEMBA1004408, HEMBA1004429, HEMBA1004499, HEMBA1004507, HEMBA1004509, HEMBA1004542. HEMBA1004596, HEMBA1004604,

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HEMBA1004776
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    HEMBA1005206,
                    HEMBA1005219,
                                   HEMBA1005274,
                                                  HEMBA1005331,
                                                                  HEMBA1005338,
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    HEMBA1005423,
                    HEMBA1005576,
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                                   HEMBA1006253,
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    HEMBA1006474,
                    HEMBA1006486,
                                   HEMBA1006492,
                                                   HEMBA1006502,
                                                                  HEMBA1006535,
                                                                                  HEMBA1006579,
     HEMBA1006648,
                    HEMBA1006659,
                                   HEMBA1006885,
                                                   HEMBA1006929,
                                                                  HEMBA1006941,
                                                                                  HEMBA1007078,
     HEMBA1007080,
                    HEMBA1007121,
                                   HEMBA1007194.
                                                  HEMBA1007300,
                                                                  HEMBA1007301,
                                                                                  HEMBA1007322.
    HEMBB1000036,
                    HEMBB1000044.
                                   HEMBB1000089,
                                                                  HEMBB1000217,
                                                  HEMBB1000215,
                                                                                  HEMBB1000272,
    HEMBB1000420,
                    HEMBB1000591,
                                   HEMBB1000593,
                                                  HEMBB1000631,
                                                                  HEMBB1000835,
                                                                                  HEMBB1000887,
    HEMBB1000908.
                    HEMBB1000975.
                                   HEMBB1000985,
                                                   HEMBB1001011,
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                                                                                  HEMBB1001112,
    HEMBB1001133,
                    HEMBB1001331,
                                   HEMBB1001337,
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                                                                  HEMBB1001367,
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    HEMBB1001394,
                    HEMBB1001429,
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    HEMBB1001753,
                    HEMBB1001797,
                                   HEMBB1001802,
                                                  HEMBB1001812,
                                                                  HEMBB1001874,
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    HEMBB1001915,
                    HEMBB1001973,
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                                                  HEMBB1001990,
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    HEMBB1002249,
                    HEMBB1002329,
                                   HEMBB1002342.
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    HEMBB1002489,
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                                                  HEMBB1002534,
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                                                                               PLACE1001983,
PLACE1002066,
                PLACE1002072,
                               PLACE1002140,
                                               PLACE1002171,
                                                                PLACE1002319,
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PLACE1002499,
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PLACE2000419,
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                                                               PLACE3000271,
                                                                               PLACE3000339,
PLACE3000341,
               PLACE3000350,
                               PLACE3000353,
                                               PLACE3000401.
                                                               PLACE4000300.
                                                                               SKNMC1000091,
THYRO1000855,
                THYRO1001559,
                                Y79AA1000065.
                                                Y79AA1000202.
                                                                Y79AA1000214.
                                                                                Y79AA1000346,
Y79AA1000784, Y79AA1000833, Y79AA1000968, Y79AA1001555, Y79AA1002220.
[0235] Clones of which expression levels decrease by TNF-\alpha are as follows:
HEMBA1002150,
               HEMBB1000240, NT2RM2000469, NT2RM2000984, NT2RM2001688,
                                                                               NT2RM4000290,
NT2RM4000496.
               NT2RM4000590.
                               NT2RM4001047.
                                               NT2RM4001582.
                                                               NT2RM4001611,
                                                                               NT2RM4001650,
NT2RM4002075.
               NT2RM4002128.
                               NT2RP1000174.
                                               NT2RP1000243,
                                                               NT2RP1000581,
                                                                               NT2RP1000688.
NT2RP1000767,
               NT2RP1000825.
                                                               NT2RP1001432,
                               NT2RP1001185,
                                               NT2RP1001286.
                                                                               NT2RP1001457,
NT2RP2000001,
               NT2RP2000248.
                               NT2RP2000841.
                                               NT2RP2001813,
                                                               NT2RP2002137,
                                                                               NT2RP2002928,
NT2RP2003517.
               NT2RP2003559,
                               NT2RP2003564,
                                               NT2RP2004933,
                                                               NT2RP2005038,
                                                                               NT2RP2006365,
NT2RP3000072,
               NT2RP3000320,
                               NT2RP3000484,
                                               NT2RP3000980,
                                                                NT2RP3001111,
                                                                               NT2RP3001420,
NT2RP3001495,
               NT2RP3002056,
                               NT2RP3002057,
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                                                                               NT2RP3002799,
NT2RP3002869,
               NT2RP3002953,
                               NT2RP3002955,
                                               NT2RP3003282,
                                                               NT2RP3003290,
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NT2RP3003385,
               NT2RP3003870,
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                                               NT2RP3004262,
                                                               NT2RP3004527,
                                                                               NT2RP4000500,
NT2RP4000524,
               NT2RP4000787,
                               NT2RP4000927,
                                               NT2RP4000955,
                                                               NT2RP4000989,
                                                                               NT2RP4001442,
NT2RP4001638,
               NT2RP4001950,
                               NT2RP4002888,
                                               NT2RP5003524,
                                                               OVARC1001270,
                                                                               PLACE1000246,
PLACE1002816.
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[0236] These are rheumatoid arthritis-associated clones.

Analysis of ultraviolet radiation damage-associated genes

- 35 [0237] It is known that ultraviolet rays give considerably adverse influence on the health. In recent years, there have been significant risks of tissue damage by ultraviolet rays because of destruction of the ozone layer. Thus, ultraviolet radiation has been recognized as a risk factor for skin diseases such as skin cancers (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with skin damage caused by ultraviolet radiation.
  - [0238] After primary cultured skin fibroblast cells were irradiated with ultraviolet ray and were cultured, a survey was performed for genes of which expression levels were varied depending on the irradiation of ultraviolet ray. First, after cultured to be confluent, the primary cultured skin fibroblast cells (Cell Applications) were exposed to  $10,000 \, \mu J/cm^2$  of 254-nm ultraviolet light.
- 45 [0239] Messenger RNAs were, then, extracted by using a FastTrack™ 2.0 mRNA Isolation kit (Invitrogen Co.) from the unexposed cells and from the cells that were exposed to the ultraviolet light and then cultured for 4 or 24 hours. The labeling of the hybridization probe was carried out by using a 1.5 μg of each mRNA in the same manner as described above. The data were obtained in triplicate (n=3). The hybridization signals were compared between the cells exposed to the ultraviolet light and the unexposed cells. The comparison was preformed by statistical treatment with two-sample t-test. Clones with significant differences in the signal distribution were selected under the condition of p<0.05. In this analysis, even when the signal is lower than others, the difference in the signal values can be detected statistically. Accordingly, clones with signal value of 40 or lower were also assessed for selection.</p>
  - [0240] Tables 353-509 show the expression of each cDNA in skin-derived fibroblast cells exposed and unexposed to ultraviolet light.
- [0241] Averaged signal values (M<sub>1</sub>, M<sub>2</sub>) and sample variances (s<sub>1</sub><sup>2</sup>, s<sub>2</sub><sup>2</sup>) were calculated for each gene in each of the cells, and then, the pooled sample variances s<sup>2</sup> were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: t=(M<sub>1</sub>-M<sub>2</sub>)/s/(1/3+1/3)<sup>1/2</sup>. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01

in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the exposed cells in comparison with that of unexposed cells.

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[0242] The expression levels of the following clones were elevated 4 or 24 hours after the ultraviolet irradiation:
                    HEMBA1001808,
     HEMBA1000542,
                                   HEMBA1002177, HEMBA1003314, MAMMA1001874, NT2RM2001100,
                                   NT2RP4000657,
     NT2RP2005732,
                    NT2RP3000592.
                                                  OVARC1000004,
                                                                  OVARC1001092,
                                                                                 OVARC1001342,
     PLACE1002816.
                    NT2RM4001002,
                                                   NT2RM4002266,
                                   NT2RM4001813,
                                                                  NT2RP2001174,
                                                                                  NT2RP2001196,
     NT2RP2005358,
                    NT2RP3000690.
                                   NT2RP3001216,
                                                  NT2RP3003464,
                                                                  PLACE1006382,
                                                                                 THYRO1000070,
    THYRO1001100, Y79AA1000342.
     [0243] The expression levels of the following clones were decreased 4 or 24 hours after the ultraviolet irradiation:
     HEMBA1000005, HEMBA1000150,
                                   HEMBA1000156,
                                                  HEMBA1000158.
                                                                  HEMBA1000168,
                                                                                 HEMBA1000231.
     HEMBA1000304, HEMBA1000307,
                                   HEMBA1000333.
                                                  HEMBA1000366,
                                                                  HEMBA1000369,
                                                                                 HEMBA1000390,
     HEMBA1000396,
                   HEMBA1000418,
                                   HEMBA1000434,
                                                  HEMBA1000464,
                                                                  HEMBA1000469.
                                                                                 HEMBA1000490,
    HEMBA1000504,
                   HEMBA1000505.
                                   HEMBA1000557,
                                                  HEMBA1000657,
                                                                  HEMBA1000673.
                                                                                 HEMBA1000682,
     HEMBA1000686.
                   HEMBA1000727,
                                   HEMBA1000752,
                                                  HEMBA1000851.
                                                                  HEMBA1000852,
                                                                                 HEMBA1000870.
     HEMBA1000872.
                    HEMBA1001085,
                                   HEMBA1001121.
                                                  HEMBA1001133.
                                                                  HEMBA1001235,
                                                                                 HEMBA1001265.
     HEMBA1001281,
                   HEMBA1001289,
                                   HEMBA1001299.
                                                  HEMBA1001303.
                                                                  HEMBA1001310.
                                                                                 HEMBA1001323,
     HEMBA1001595,
                   HEMBA1001620.
                                   HEMBA1001640.
                                                  HEMBA1001678.
                                                                  HEMBA1001712,
                                                                                 HEMBA1001835.
    HEMBA1001950.
                   HEMBA1001987.
                                   HEMBA1002253.
                                                  HEMBA1002321,
                                                                  HEMBA1002341,
                                                                                 HEMBA1002419,
                                   HEMBA1002818,
    HEMBA1002679,
                   HEMBA1002728,
                                                                  HEMBA1002999,
                                                  HEMBA1002935,
                                                                                 HEMBA1003034,
    HEMBA1003071,
                   HEMBA1003098.
                                   HEMBA1003142,
                                                  HEMBA1003175,
                                                                  HEMBA1003202,
                                                                                 HEMBA1003212,
    HEMBA1003220,
                   HEMBA1003276,
                                   HEMBA1003373.
                                                  HEMBA1003417,
                                                                                 HEMBA1003528,
                                                                  HEMBA1003447,
    HEMBA1003684,
                   HEMBA1003799,
                                   HEMBA1003885,
                                                  HEMBA1003989,
                                                                  HEMBA1004011,
                                                                                 HEMBA1004055,
    HEMBA1004133,
                   HEMBA1004225,
                                   HEMBA1004272,
                                                  HEMBA1004353,
                                                                  HEMBA1004631.
                                                                                 HEMBA1004669.
    HEMBA1004705,
                   HEMBA1004753,
                                   HEMBA1004776,
                                                  HEMBA1004803,
                                                                  HEMBA1004816,
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    HEMBA1005047,
                   HEMBA1005079,
                                   HEMBA1005101,
                                                  HEMBA1005149,
                                                                  HEMBA1005152,
                                                                                 HEMBA1005202
    HEMBA1005314,
                    HEMBA1005372,
                                   HEMBA1005511,
                                                  HEMBA1005513,
                                                                  HEMBA1005518,
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                                                  HEMBA1005609,
                                                                  HEMBA1005632,
                                                                                 HEMBA1005853,
    HEMBA1006031,
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                                                                                 HEMBA1006696,
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                                                                  HEMBB1000399.
                                                                                 HEMBB1000434,
    HEMBB1000438,
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                                   HEMBB1000673.
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                                                                                 HEMBB1000883.
                                   HEMBB1001182,
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	[0244] These clo	nes are ultraviolet r	adiation damage-as	sociated clones.		

Table 12

Expression of each cDNA in human tissues (containing clones that are not described in Examples.)

	Clone_name	Hoars	Lung	P. gland	Thyous	Brain	Ridnoy	Livor	Sploon
	GAPOH (Cr1)	38.210	32.670		13.580	11.230	21.120	24.910	22.440
	Bactin(Cr2)	279.280	368.870	111.100 1	17.500	92.880	114.650	82.990	256.790
	ADRGL 1000005	53.882	23.005	32.749	22.858	26.564	24.940	22.644	27.001
5	ADRGL 1000007	94.778	85. 185	160. 457	57.191	101.768	62.489	67.150	73. 543
	ADRGL 1000009	11. 141	50. 520.	10. 357	7.177	6.013	5. 219	14. 272	21. 225
	ADRGL 1000011	71.656	24. 579	29. 358	19.473	24. 898	30.747	49. 220	22.221
	ADRGL 1000027	36. 238	25. 252	20. 855	7.328	11, 196	14. 298	19.658	11. 288
	ADRGL 1000058	66. 209	129.497	55. 226	49. 241	30.219	55.872	67.027	243. 436
10	ADRGL1000069	38.630	23. 459	28. 991	12.540	27.353	33.633	28.774 67.906	20.911 89.439
, 0	ADRGL 1000077	97. 465	63. 656	448. 427	83.412	71.108	53. 740 73. 339	96.037	73.091
	ADRGL 1000092	89. 423	45. 692	55.810	26.033	44.148 25.964	41. 391	42.837	29.666
	ADRGL 1000099	73.675	63.974	36. 128 77. 017	17.024 24.777	33.549	58. 986	295.009	84. 985
	ADRGL 1000 136	141.745 394.563	155. 829	271.210	92.899	165. 627	251.266	253.420	150. 294
	ADRGL1000159	50.073	25. 425	39. 296	15. 194	16. 125	20.040	33.720	23. 278
15	ADRGL1000160	69. 386	31.051	59. 416	20.154	39.799	27.027	47.169	20.716
	ADRGL 1000171	57.047	23.011	43.063	23.860	40.581	59.814	117.055	32.630
	ADRGL 1000 181	45. 892	18.666	34. 476	15.434	34. 225	32. 962	39.693	16.334
	BGG111000015	153. 242	42. 337	92.865	41.003	45.168	88. 524	85.990	73.392
	BGG111000016	177. 367	94.731	119.688	34.159	30. 249	98.806	98.783	39. 204
20	BGG111000017	84.712	32.614	38. 131	20.878	18.769	32.340	39.666	20.750 11.937
	BGG111000022	52.468	20. 452	67. 167	12. 167 12. 585	11.158	18. 241 15. 525	19. 197 16. 757	13. 406
	BGG111000031	30.008	17. 072 36. 336	40.883 51.176	26.964	43. 122	43.770	49, 107	38.776
	8GG111000042 8GG111000046	49. 926 31. 618	26. 472	34. 182	31.854	12.650	25.784	18. 430	25, 385
	BNGH41000020	5031. 103	2993. 496		537.162				3649.144
25	BNGH41000025	91.717	35. 026	73.901	27.713	30.765	36. 523	37.596	47.074
25	BNGH41000026	176.757	77. 439	98. 345	35. 807	56.991	91.310	75.797	70.241
	BMCH41000027	65.029	56.353	25.896	22.494	12.763	23.748	17.836	23.859
	BMGH41000035	148.779	66.776	119.727	56.576	60.996	96.959	72.461	64.458
	BMGH41000037	79. 500	29.611	43. 438	18.317	20.857	36.272 125.323	27.525 86.783	122.959
	BNGH41000042	224. 484	110.084 32.253	168.448 54.063	104.351 14.729	102.259	22. 435	29.566	28. 937
30	BNGH41000048 BNGH41000056	56. 144 67. 258	18.694	30.075	15.602	10.072	20.735	16.100	7.642
	BNCH41000087	98. 262	46. 173	77.657	35. 329	40. 900	50.029	50.841	45. 285
	BNCH41000091	50.895	16.985	28.392	10.147	5.469	22.794	10.725	12.410
	BNCH41000157	69.043	34.730	40. 597	18.088	27.072	22.074	25. 410	24.950
	BMGH41000169	44.850	21.770	28.655	11.403	25. 991	28.509	25, 634	25.843
35	BNCH41000181	17. 163	15.689	13.948	3.996	9. 287	13.139	15.553	16.575
	BMGH41000198	81, 510	36.250	60.860	20. 585	26.929	35. 751 15. 235	31.695 27.285	28.325 35.709
	BNCH41000219	30. 302	25. 156 65. 948	22. 187 93. 499	13.757 51.108	11.208 92.555	101.245	96.716	78. 266
	BNGH41000229 BNGH41000237	252.790 85.757	46, 997	55. 170	26.780	33.764	47.456	37.007	39.131
	BNGH41000238	17.744	36. 938	42.360	14.922	35.749	42.848	39.238	13. 241
40	BNGH41000243	45. 446	23.667	44. 798	20.875	10.516	23.918	22.443	27.033
70	BNGH41000270	60.889	18.651	29.618	10.724	15.979	12.351	19. 152	22.314
	BRAUH100004	43.673	28.539	7.640	11.388	19.198	14.903	32.353	23.777
	BRAGH1000018	59.409	17. 941	102.270	17. 107	709.078	25.732	24. 214 66. 754	24.767
	BRACH1000021	104.772	29. 951	67. 089	32.199	1169.154 64.521	55.762 70.731	79.670	40. 928
	BRAUH1000027	152.205	47.310	55. 840	40.856	59.552	56. 487	64.886	100.132
45	BRAUH1000029	29, 419	16.761	31.101	16.622	30, 633	18. 200	17.998	15. 196
	BRACH1000050	161.264	71.786	118.976	51.863	61.542	97.720	81.271	69. 194
	BRACH1000051	74.067	34. 341	44,047	20.726	30.434	42.055	53, 856	24.624
	BRAUH1000050	68.789	22.598	35.012	16.493	19, 127	38.662	34. 923	28.094
	BRACH1000075	17.318	16, 898		8.901	18. 133	17.219	9. 321	11. 200
50	8RADH1000081	43.025	12. 998		7.655	123.677	17.673	15.924	9.844
	BRAUH1000084	174.384	42.178		47.752	152, 188	77.111	746.700	102.296 35.985
	BRAUH1000095	118. 239	59.676		28.174		52.880	52.427	
	BRADH1000096 BRADH1000097	95, 841	44. 967 72. 506		65.637	64.200	73.707	63.827	63.762
	BRACH1000100	11.943	19, 037		13.536		16.582	16.646	
55	BRAUH1000101	134.838	57. 232		40.741		71.642	88. 432	57.336
30									

Table 13

	8RAUH1000104	25.414	18, 303	14.825	7.695	38, 918	23, 970	23.794	11.048
	88AUH1000107	16.949	5.616	12. 463	5.518	6. 355	5. 084	9, 107	6.573
	BRADH1000110	615, 476	492.704		383.612	368, 156	369.621	277.348	340. 450
5	BRADH1000111	175. 556	68. 459	92.209	45.974	64.703	81.723	90.369	57, 301
	BRADH1000135	199, 303	38. 098	72.093	26.809	57.720	91.668	87.016	35.866
	BRAUH1000190	56.386	41.640	57.914	22.782	55.671	40, 034	35, 280	40, 134
	HEHBA1000005	11.985	23. 427	18.882	9.766	12.656	9, 959	23. 443	21.677
	HEMBA1000006	37. 398	24. 521	24. 529	15. 587	22.317	13. 336	16.038	15, 295
	HEMBA1000012	81.820	57. 193	66.828	26.683	55. 423	58. 731	85.614	66.259
10	HE23BA 1000020	157.967	64. 157	115.635	51.940	77. 293	77.321	83. 989	74. 362
	HELIBA 1000030	82.882	35. 447	66.058	26.464	40.990	60.871	47.058	50.652
	HEMBA1000034	47.434	17.878	50.696	5. 594	14.005	5, 573	24.652	7.134
	HEMBA 1000042	147.376	94.003	330.908	69.071	76.472	55. 477	37.783	60.479
	HEMBA1000045	28. 478	20. 289	20. 548	12.445	11.835	22.788	11.196	15.775
	HEMBA1000046	85.160	84. 475	242, 940	57.017	68, 488	45. 288	37.098	47.486
15	HEL'BA1000047	21.380	18.899	18.166	11.393	11.185	12. 292	6, 491	12.018
	HEMBA1000048	243.559	55, 114	84. 448	24.247	43.131	99.333	57.041	37. 362
	HEMBA 1000050	22.711	11.876	21.972	7.477	4.096	13.675	10.347	7.770
	HEMBA1000053	45.071	26.410	38.158	15. 982	30.754	36.740	34. 184	24. 269
	HEL'8A1000060	101, 197	34.766	50.643	19.938	34.641	54.061	42.309	22.530
	HEMBA1000072	240.166	213.938	224.688	163.030	115.246	207.809	112.361	276.098
20	HEMBA1000073	23. 202	9. 580	10.815	1.698	6.680	18. 155	12.304	14. 973
	HEL'BA 1000076	95.997	46.783	177.931	32.617	48.964	50.792	33.947	44, 142
	HEMBA1000084	66.603	25.710	48. 434	18.006	22. 553	38.118	40.479	29.683
	HEMBA1000087	70.084	17.515	26.544	8.450	17. 590	29, 220	19.519	22. 565
	HELBA1000088	15.474	8.614	19.903	4,775	4.519	11.446	34. 905	6. 528
2.7	HEL'8A1000091	80.622	38.604	59.393	23. 956	44. 939	49.760	33.946	24.614
25	HEMBA1000111	85.814	95.270	270.642	75. 147	54. 384	70.071	29. 529	55. 422
	HEMBA1000121	55.476	43.368	146.465	37.419	29. 398	30.694	17.702	30. 398
	HEMBA1000128	37.278	27.165	34.516	13.619	17.702	28.069	12.834	23. 965
	HEI:BA1000129	51.488	19.659	44. 907	12. 208	27.243	30.959	24.383 8.907	26.851 20.635
	HEE:3A1000141	12.961	24. 515	32.107 20.418	14. 353 8. 202	13.502 9.575	11. 152 14. 877	10.000	7.817
30	HEMBA 1000146 HEMBA 1000150	29. 273 534. 562	11.479 326.814	684.147	211.774	218. 448	322. 246	235.752	256.883
30	HEMBA1000154	95. 272	92. 253	101.483	54. 276	42.896	75. 526	92.689	188.019
	HEMBA1000155	50.177	72. 591	58.026	31.149	21.865	38. 964	27.634	50. 220
	HEMBA 1000158	260.718	63. 920	89.680	36.337	44.915	93. 421	111.344	53. 562
	HEMBA 1000168	74.416	61.152	62.826	30.512	23. 287	34. 966	44.005	33.564
	HEMBA 1000180	28, 502	22.412	28.571	11.701	19. 230	10. 903	11.731	14, 102
35	HEMBA1000185	115.723	50.661	213.994	51.166	43.435	56. 261	38.862	44. 992
-	HEMBA1000188	21.302	14.879	16.948	11.392	11.821	10.656	12.501	6.979
	HEMBA 1000193	14. 122	8.318	11.905	7.519	4.736	3. 349	8.544	7.842
	HEMBA1000194	54.688	49. 534	143.817	37.736	20. 221	34. 328	23.359	56.497
	HEMBA1000201	21.062	14.098	8.690	6.237	5. 109	5. 059	9.317	10.522
	HEMBA1000213	22. 388	25. 532	25.777	8.470	17.320	9. 084	8.469	11.766
40	HEMBA 1000216	65. 935	51.368	92.680	19. 202	33.659	40. 971	36.328	34.891
	HEMBA 1000227	52.577	31. 332	34.925	19.503	18.411	21.504	22.590	25. 781
	HEMBA1000231	114.369	54. 299	131.256	38.550	43.246 93.250	29.778	24. 266 49. 319	30.410 59.288
	HELBA1000237	91.024	91.360	199.338	58, 292	25, 396	57.000 32.604	38.910	32. 153
	HEMBA1000243	53.456 173.469	43.969 104.733	111.513	33.079	65. 527	124, 532	90.927	78.610
	HEMBA 1000251	22.709	12. 333	14. 367	9.019	16.095	13. 221	11.516	11.018
45	HEMBA1000254	74.060	35.626	130.009	20.848	37.481	24. 002	20.553	13.215
	HETBA1000264	29. 478	15. 248	23.537	9.473	3.863	11. 228	13.690	3.797
	HEP3A1000269	36.718	13.465	28.932	20. 412	9.705	12.833	7.348	24. 793
	HEMBA1000275	66.201	39.367	84.077	38. 846	77.871	49. 267	36.211	38, 871
	HEMBA1000280	33.299	36.073	54.357	24.720	38, 017	35.751	21.696	30.785
	HEMBA1000282	93.815	121.083	171.037	93.484	123. 971	70.384	56.916	92.414
50	HEMBA1000287	12.439	24. 935	29.793	10.840	37. 925	9.632	2.866	7.311
	HEMBA1000288	45.269	30.009	145.363	25.471	9.769	16.272	9.701	15.510
	HEMBA1000290	14.803	5.750	10.615	5.725	2.559	8.602	8. 358	9. 224
	HEMBA 1000296	27.085	22.625	21.195	9.790	16.909	12.402	15. 289	17. 159
	HEMBA1000300	98.491	119.119	304.884	73.660	85. 595	48. 175	43.495	66.547
55	HEMBA1000302	23.840	15.442	27.722	16.143	13.081	13.879	8. 259	12.569
55						-			

Table 14

	HEHBA1000303	129. 286	51.013	88.777	32.513	50. 462	82.994	44.818	49.271
	HEHBA1000304	112.022	67.470	328.677	54.678	79. 305	43. 526	38. 469	55. 762
	HEMBA1000307	14.054	22.013	31.964	13, 167	15.571	7.974	10.014	8.685
5	HEMBA1000312	97.082	69.330	183, 923	45. 322	45.087	52, 968	37.741	38. 246
	HEBBA1000318	15.164	16. 264	18.766	11.688	3.620	10.732	8.295	14.675
	HEMBA1000327	29.404	59.618	81.347	41.731	85.004	48. 526	49. 421	46.866
	HEHBA1000333	16.964	13.930	14, 530	1.872	5.778	1.571	0.392	3.743
	HEMBA1000338	121.878	62.572	348. 751	55.463	49.114	38, 561	30,698	40.644
	HEHBA1000343	25. 229	29. 781	46. 395	20.673	5.872	16. 551	10.139	14. 088
10	HEMBA 1000349	23.061	12.586	31.755	7.020	17.658	11. 522-	14.807	15.611
	HEMBA 1000351	92.847	57. 338	196. 577	41.762	37.094	35. 370	27.645	28.615
	HEMBA1000355	85. 210	38. 388	64, 299	18. 101	33.114	43.511	37.808	26.628
	HEBBA1000356	60.438	38.786	62.442	20.784	17.694	38.058	40. 431	28.899
	HEMBA1000357	84.898	55.990	206.803	54. 151	42.793	39. 432	26.076	44. 579
	HEMBA1000356	47.131	42.031	90. 450	27.056	20.718	23.499	14.632	23. 547
15	HEMBA1000369	71.428	40.685	54. 384	17.613	21.422	34. 985	37.622	36.900
	HEEBA1000370	16.354	14.949	22.988	7.916	18.390	15.359	13.426	6.647
	HEMBA1000376	80.183	75. 300	201.705	55. 266	66.687	44. 612	55, 386	56.070
	HEEBA1000387	100.497	129. 367,	351.196	80.257	104. 250	74.007	57.619	79.876
	HEMBA1000389	69.342	34.021	71.118	22.346	27.319	47.935	53.026	34. 161
	HEMBA1000390	19. 206	25.788	21.028	12.401	18.372	13.751	16.243	15.036
20	HEMBA 1000392	19.400	22.884	44. 179	8.776	11.742	10.594	12.266	12.463
	HEMBA 1000396	75.409	50.195	81.870	27.979	30.393	31, 235	17.771	19.584
	HEMBA1000411	35. 966	24.397	25.987	10.341	31.398	31. 214	50.056	18.580
	HEMBA1000418	8.165	10.778	14.987	4.031	12.495	7.913	6.363	2.306
	HEMBA1000422	93.699	38. 329	85.266	39.826	45. 992	44. 729	42.886	34. 308
0.5	HEMBA1000428	51.017	30.690	79.229	26.579	24.840	17. 767	18. 424 2. 788	18.608 2.756
25	HELBY 1000434	1.747	3.214	11.346	1.210	1.602	2. 927 17. 969	11.723	10.645
	HEMBA1000442	21.750	7.698	16.227	7. 252	3.336 16.860	31.691	47.856	102. 287
	HEMBA 1000443	67.291	35.910	34.775	26. 420	34. 107	119.377	108.645	60.266
	HELBA1000445	236. 986	69.546 37.560	90. 283 63. 451	32.233 22.640	41.092	65. 256	62.972	43.493
	HEL'BA1000456	95.368	35.333	74.945	20.475	25.324	26. 253	13.654	31.317
30	HEMBA 1 000459 HEMBA 1 000450	18. 649	27.246	21.973	9.613	15. 230	14.091	9.746	16.955
30	HEP-8A 1000452	220. 184	42.636	96.490	31.332	83.626	109.503	92.971	62. 126
	HELTBA 1000464	34.277	15. 137	27.210	10.862	15.595	20.793	16.716	16.539
	HEM8A1000468	41.755	41.852	68.356	10.400	23.452	43.909	24.048	22.968
	HELBA1000469	68.229	71.011	256.705	47.535	29.853	34. 188	22, 568	39. 190
	HEPBA1000477	185. 220	47,546	102.939	26.276	40.188	95. 247	52.454	28. 109
35	HEMBA1000481	47.276	37.528	24.407	17.115	24. 182	29.826	20.717	25.819
•	HEL'BA 1000488	95.226	31.249	71.522	21.667	27.715	44. 499	53.708	33.306
	HEYBA1000490	29.915	13.747	32.568	14.002	12.056	6.900	11.274	7.559
	HEP3A1000491	80.198	22.903	47.786	20.675	32.551	52.682	37.109	28. 282
	HEMBA1000498	191.186	112.767	454.998	88.614	102.997	82.927	53.205	120.837
	HEMBA 1000501	57.318	55, 923	180. 158	44. 170	27. 291	34, 954	18.532	34. 117
40	HEYBA 1000504	1.033	5.893	7.152	1,726	0.520	2. 245	2. 551 19. 508	1.091 31.925
	HEMBA 1000505	55.746	36.631	48. 155	21. 562	14.691	97. 289	230, 331	95. 150
	HEMBA 1000507	204. 165	114.530	305, 249 309, 791	86. 138 72. 709	81.505 70.180	77.388	63.849	45. 940
	HEMBA 1000508	205.724	105.067	31.505	16.650	14.796	15.847	24.729	17.601
	HEMBA 1000518 HEMBA 1000519	39. 157 166. 937	29.100 142.675	468, 435	148.478	123.978	128.646	85.670	111.078
	HELBA1000520		0.000	0.000		10.619	1.488	9.513	9.395
45	HEPBA1000523		22.090	40.875	13.852	21.603	32. 384	20.478	21. 422
	HELBA1000531		34.044	40.027	12.264	11.034	29.775	20.421	12.540
	HEMBA 1000534		0,000	0.000		48.940	25. 365	41.242	72. 583
	HEMBA 1000538		0.000	0.000		19.981	17. 606	26.698	23. 904
	HEMBA1000540		47. 343	33.145		27.059	33. 931	15.639	31.893
	HELBA1000542		33. 152	58.093		35.278	55, 508	47.917	47.623
50	HEYBA1000545		136, 401	48.802		12.534	7, 119	25.484	15.094
	HEPBA 1000547		20.199	32.694		22. 359	12.020	13.535	20. 227
	HEMBA1000551		171.089	543.876	131.764	115.775	116.646	69.596	152.516
	HEMBA1000555	10. 531	20. 199	25. 801	24. 488	14.071	15. 431	5. 986	10. 933
	HEMBA 1000557		48. 396	168.724			31.872		30. 926
	HEMBA 1000561	56. 992	22.797	51.047	10. 187	16.301	34.904	24.661	22.470
55									

Table 15 ·

	HEMBA1000563	9.473	11 545	19 205	6 120	12 600	10 122	2 010	14 002
			11.545	18.205	6.139	12.689	10.132	7.939	14. 253
	HEMBA1000567	41.385	38.483	27.881	32. 207	15.544	26.052	15.086	86.601
5	HEMBA1000568	44.686	33.379	126.524	26.300	22.533	17.402	26.970	18.707
· ·	HEMBA1000569	58. 184	27. 187	41.012	21.787	12.925	36.191	33.944	23. 225
	HEMBA1000575	155,833	155.759	434.526	92.140	79.143	69.949	59. 928	71.189
	HEMBA1000588	41.087	26.072	31.610	14. 580	18.024	18, 458		
	HEMBA1000590							23. 553	13. 279
		29.693	17.090	23.618	7.069	6.633	16.725	20.068	13.042
	HEMBA1000591	106.772	54.874	98.079	34.099	31.776	57.170	48. 488	32.766
10	HEMBA1000592	7. 408	10.031	9.435	9. 551	8.209	5. 142	7.480	10.319
	HEMBA1000594	18.401	11.048	22.547	15. 327	9. 596	12.099	8.751	6.852
	HEMBA1000604	96.047	78.462	146.030	49.571	36.099	70.815	41, 797	47, 748
	HEMBA1000607	46.819	15.606	46.037	9. 438	19.149	21.038	17.317	25, 404
	HEMBA1000608	8, 985		6,705					
			3.040		0.000	7. 378	4. 453	0.000	5. 544
	HEMBA1000622	45. 570	55.746	113.666	40.310	18. 167	19.390	15, 895	29. 149
15	HELBA1000634	126.532	49.146	138.073	29.094	95.787	79.662	60.271	71.657
	HEMBA 1000636	151.899	51.270	126.200	39.161	51.864	62.611	54.056	39. 415
	HEMBA1000637	33.241	23.587	39.380	18.047	16.265	30.075	28. 226	24. 559
	HEMBA1000655	80, 165	70.766	219.283	58.901	61.320	45. 821	40.741	62. 639
	HEMBA1000657	60.961	31.993	41.401	18.008	30.565	35. 201	35.611	42. 178
	HEMBA1000662	8,600	8. 490	11.263	5. 475	2.201			
	HEMBA1000664						6.140	1.557	2. 504
20	HEMBA1000671	14. 358	5. 082	3.637	2.670	3.516	4.913	3.094	3. 579
		11.588	15. 473	26.067	17.940	8.865	7.647	10.779	21.196
	HEMBA1000673	73.174	77.410	193. 253	46.051	34. 388	33. 975	25.896	31.646
	HEMBA1000675	7.666	12.047	22. 123	5.764	42.036	15.788	10.254	15. 555
	HEMBA1000678	7.453	12.314	21.083	12.174	14.897	12.628	6.969	6. 584
	HEMBA1000682	118.965	125.696	255.731	86.894	61,443	66.299	49,060	82.939
25	HEMBA1000686	25.079	17. 463	23.126	12.722	10.282	13.835	21. 393	18. 154
25	HEMBA1000702	206.683	94. 357	266.585	62.386	79.930	90.914	98. 397	60. 559
	HEM8A1000705	25. 430	25.862	47.190	13.191	19, 599	26, 364	25,013	18. 833
	HEMBA1000713	56. 893	25. 288	70.751					
	HEMBA1000718	50.149			17.660	24. 138	23, 311	21.805	21.736
			43.869	128.515	28. 289	23. 213	18. 458	10.003	17.419
	HELBA1000719	37.969	17.467	28.513	12.147	12. 768	22.543	14.744	14. 432
30	HEMBA1000722	15, 150	9.762	14.699	6.768	11.726	12.080	5. 907	9.953
	HEM8A1000726	159.817	111.276	463.937	91.448	109.093	58. 587	46.517	70.087
	HEMBA1000727	22.867	26.803	28.886	21.475	11.199	14.966	8.634	30. 401
	HEMBA1000732	28.630	11.011	12.790	4.617	3.548	13.325	19.978	13.472
	HEM8A1000736	24. 568	21.982	21.410	7.431	11.376	41.026	31.698	16.801
	HEMBA1000743	0.741	4. 467	1.793	1.637	1.227	3.642	4. 563	3. 368
	HEMBA1000745	8. 930	7.067	14. 546					
35	HEMBA1000747	21.442			3.314	10.067	5. 403	9. 225	6.085
	HEMBA1000748		12.487	25.662	17.081	5. 384	10. 287	9.865	8. 267
		22.924	14. 885	35. 721	12.634	3.045	11.508	4.110	11.756
	HEM8A1000749	67. 267	50.826	159.211	43.879	20. 345	29.613	19.447	31.693
	HEMBA1000752	54. 929	35.778	162.005	28. 209	31.540	25. 132	15.650	20.776
	HEMBA1000753	120.889	83. 878	155.892	48.092	54.307	53. 238	38. 941	39. 331
	HEMBA1000757	20. 234	22. 592	52.608	29.935	23,071	24. 503	14.548	43.779
40	HEMBA1000760	12, 599	38.665	19.973	15.800	30, 188	14, 155	10.570	39. 229
	HELBA1000769	114.956	74. 924	304.424	66.815	39. 365	48. 405	39.918	55, 931
	HEMBA1000773	2.162	5. 360	11.883	4. 445	0. 965	3. 158	3. 956	2. 663
	HEL'8A1000774	128. 563	115.732	330.111	84.461	69. 618	59. 363	42.656	
	HEMBA1000780	5. 850			6. 924				56. 152
			7. 130	24.176		6. 903	6. 546	6.667	9. 576
45	HEMBA1000783	8. 127	5. 076	13.701	3. 276	8. 863	6. 241	5. 435	4. 429
70	HEMBA1000791	41. 433	51.546	108.542	29.633	42. 735	44. 515	43. 187	40. 856
	HEMBA1000793	108.761	30.885	54. 568	18.670	31.512	54.669	45.458	34. 788
	HEMBA1000802	15.062	11. 125	9.052	10.300	11.505	12, 950	15.354	16.952
	HEMBA1000813	106.763	52.683	69.701	32,507	44. 369	65.862	59.842	56.799
	HEL'8A1000817	19, 480	7.070	17.915	4.016	15. 239	18. 434	11.273	8.079
	HEPBA1000822	9. 520	10.358	15.760	7, 218	8.704	11. 185	6.639	4. 662
50	HEMBA1000827	96.001							
			12. 420	24. 041	8. 305	24.000	6.709	3. 488	8. 591
	KEMBA1000833	53.675	28.970	35.897	14.604	26. 383	29.035	20. 591	14, 341
	HEMBA1000835	74.696	67.353	83.737	34, 349	42. 834	61.145	66.784	52.015
	HEMBA1000843	74.227	54. 197	92.042	37.825	58. 573	98. 943	87.569	55. 077
	HEMBA1000851	23.913	14.070	13.081	6.847	8. 634	12.419	19.200	22, 286
	HEMBA1000852	56.702	54.074	105.085	31, 127	34. 200	31, 843	28.843	30. 311
55	HEMBA1000867	15. 548	10.247	11.912	6.256	1. 227	12.374	8.518	5. 611
							,	0.310	<u> </u>

Table 16

	EIPHOALAGOOAA I	10 700 1	10 205 1	24 020 1	15 AC1	6 020	12 200	14. 305	14.712
	HEMBA1000869	19.696	18. 785	34. 039	15.061	6. 930	13. 298		17. 508
	HEMBA1000870	64. 189	38. 245	44.665	12.647	23.970	41.195	21. 911	
5	HEMBA1000872	46. 848	46.546	86.933	36.087	40.608	42.532	43.479	36.141
•	HEMBA1000875	35. 460	41.165	32.238	11. 297	35. 077	29.781	19.453	23. 540
	HEMBA1000876	89. 976	56.654	194.869	42.595	57.670	53.567	36.331	40.884
	HEMBA1000907	22. 959	9,656	10. 917	3.599	3. 363	5. 327	13.032	10.676
	HEMBA1000908	45, 409	18. 456	30.665	12.448	8, 174	19.529	24.789	16.299
	HEMBA1000910	47, 107	13, 681	26.933	5. 866	7,073	19.938	22.971	11.592
			29. 880	114. 873	25. 206	16.670	25.895	26.769	24.710
10	HEMBA1000918	67.437				16.008	24.639	23.073	20. 233
	HEMBA1000919	44. 938	29.704	40. 184	22.126		53. 492	33.824	20.798
	HEMBA1000934	162. 546	35. 314	59.012	18.820	30.796			
	HEMBA1000935	16. 284	29. 481	71.669	12.587	23.834	13.188	7.830	13.322
	HEMBA 1000940	44. 243	39. 296	75.619	25.080	28, 113	39. 401	25. 948	30. 168
	HEMBA 1 000942	126.095	96.812	260. 912	62.657	49.118	47. 891	35.814	49.631
15	HEMBA1000943	14. 439	12.702	14, 690	4.792	8. 391	11.856	11.039	7.414
13	HEMBA1000946	15.461	5. 506	18.692	9,000	5.772	0.000	19.405	9.939
	HEMBA1000960	179.860	151.073	343.747	107.319	85, 691	117.093	82.928	94. 494
	HEMBA1000962	73. 395	34, 803	60.061	26.562	28. 789	47.944	60.067	31.619
	HEMBA 1000968	14, 529	12. 486;	35, 270	18.733	6.213	7. 458	7.214	4.624
	HEMBA1000971	50, 148	19. 281	37.515	12.222	19, 562	29.874	22.045	23.135
	HEMBA1000972	51, 100	33. 450	188, 137	28.972	24, 576	23.736	13.731	27.272
20	HEMBA1000974	5. 609	10.649	12.866	2.929	2.603	3.800	6, 104	4. 964
		34. 417	19, 132	42, 499	15.644	4, 009	16. 478	14. 192	14. 353
	HEMBA1000975			99.641	19.754	27.516	38. 801	31. 347	36.440
	HEMBA1000979	90.061	38. 532		19.544	34. 385	38. 280	24. 897	29.059
	HEMBA1000981	35.338	31. 281	38.672			32.384	27.465	31.286
	HEMBA1000983	71.391	34. 501	58.683	22.640	32. 825			
25	HEMBA1000985	9. 290	20. 363	22.497	4.058	6.343	9. 035	7.852	3. 257
20	HEMBA1000986	128.714	74.713	236.019	56.662	52. 957	85. 340	63.718	54.892
	HEMBA 1000991	72.707	55. 780	160.717	34.676	32. 494	41.317	23.483	37.846
	HEE:8A1001007	123.690	42. 563	69.807	23.525	34. 263	47.777	47.496	48. 154
	HEMBA1001008	124.864	47.842	83.746	18. 125	25. 490	52.693	30.668	24.961
	HEMBA1001009	-37.843	29, 269	36.715	11.055	17.115	17. 937	17.701	22.055
	HEMBA1001014	109.049	83, 356	233. 234	60.123	61.977	94. 424	47.095	74.625
30	HEMBATODIUTY	50.408	20. 212	48.394	16.020	28. 537	31.917	27.876	24. 283
	HEMBA1001019	7.327	7. 582	14.865	6, 154	10.598	5. 643	3. 920	7.188
	HEMBA1001020	53.067	55. 646	115.814	31.640	25. 647	24. 596	23.146	27.169
	HEMBA1001021	115.724	42.415	59.434	28.828	26. 181	64.484	64.173	29.614
	HEMBA1001022	37.883	25. 835	28.969	18.452	20.270	22.790	25.194	20.783
	HENBA1001024	23. 524	15. 235	16.511	8, 023	11,818	13.894	8.606	8.098
35	HEMBA1001024	21.343	12.515	18.851	6.888	7, 288	12.663	8.419	7.418
				9, 892	10.750	19, 163	9. 299	8.047	8. 589
	HEMBA1001043	10.374	11. 995		100. 376	67.510	61.660	46.295	68.994
	HEMBA1001051	124. 869	115. 181	387.345		11.445	24. 382	15.726	12.323
	HEMBA1001052	38.892	13.860	19.067	12.855	26, 220	46, 725	42.356	36. 506
	HEM3A1001059	98.097	41. 525	66.565	27.826				52.612
40	HELBA1001060	116.857	74.020	161.485	61.750	50.524	52. 957	38.575 17.387	16.720
40	HEMBA1001064	32. 251	24. 026	33.937	14.007	7.907	13.710	6.779	24. 242
	HEMBA1001071	25.850	16.043	19.924	7.855	3, 425	9, 530		
	HEMBA1001077	24.689	23.055	64.486	19.413	16.821	16.858	13. 165	12.873
	HEMBA1001078	33.254	26.761	41.713	26.498	24. 531	31.498	25. 302	23.636
	HEMBA1001080	57.701	23. 951	31.254	22. 489	24. 848	33. 265	31.880	26.484
	HEPBA1001084	62.698	41.625	171.096	31.438	31.760	24. 829	17. 487	26.581
45	HEMBA1001085	159. 252	116.909	294.247	77. 235	81.384	76.498	59. 989	55.574
	HEMBA1001088	74.704	42.537	46.695	19.266	25. 146	33.498	44.927	26.310
	HEMBA1001093	30.048	28.810	72.081	20.831	14.610	11.033	15.558	22. 531
	HEMBA1001094	5. 535	8.779	10.059	3,089	4. 628	4. 521	4.834	4. 468
	HEMBA1001099	18. 322	24. 021	14.814	7.146	13.778	16.055	11.044	10.190
	HEMBA1001104	21.919	13.788	35.048	9.637	18.058	24. 450	21.559	18.527
50	HEMBA1001109	186.384	190. 240	540.908	155. 496	134, 530	93.324	78.690	116.187
50	TIET DAILOUING			187.547	75.857	35, 109	66. 259	69.432	341.702
	HEMBATCH1114	89.023	252. 529				18. 209	9.519	15.621
	HEMBA1001121	32.820	25. 812	89.860	19.710	34. 244			
	HEMBA1001122	3.304	6.213	8.316	4.763	19.120	5.650	4. 506	23.059
	HEMBA1001123	108.859	55.807	190.789	41.415	39.028	42.683	25. 551	30.174
	HEMBA1001133	50.744		36.786	14.764	34.752	26.702	23. 524	11.367
55	HEMBA1001137	38.685	21.659	46.297	21.567	13.174	15.867	11.767	25. 508

Table 17

	UCUDA 1001140	CO 450	CC 160 T	100 101	40 027 1	60 262 1	44 402 [	20 267	43.561
	HELIBA1001140	60. 453	66. 122	169.353	48.837	60.363	44.403	30.367	
	HEMBA1001144	278.125	195.811		207. 291	166.089	101, 134	106.337.	142. 120
5	HEMBA1001145	58.539	241.358	206.084	46. 342	39. 316	61.827	91.170	66.852
	HEMBA1001158	29.417	28. 121	43.877	13.337	24. 176	19. 965	18.089	28.622
	HEMBA1001172	74.727	47.695	213.708	37.115	24.460	26. 520	19.178	32.709
	HEMBA1001174	6.279	8.617	8.831	7.914	2.574	8.031	3, 119	4. 980
	HEMBA1001175	29.561	34.909	43.568	19.819	34.829	16.588	19.883	17.824
	HEMBA1001182	136.762	64, 608	105, 979	44. 066	83.417	86.736	126.297	79.785
								6.347	
10	HEMBA1001184	16.758	9.703	22.060	9.016	11.018	10.205		9. 176
	HEMBA1001192	15.119	10.798	11.626	6.559	5. 736	3, 435	9.089	11.273
	HEMBA1001197	82.571	114.743	110.687	83. 431	56.396	68. 797	99, 959	173. 379
	HEMBA1001208	40.250	30.964	37.220	19.514	11.451	24. 172	27.637	12.469
	HEMBA1001213	81.501	37.345	57.618	18.958	24.480	52, 160	51.978	31.326
	HEMBA1001214	36.798	16.011	20.958	17.612	12.418	20.697	19.108	21.328
	HEMBA1001221	14. 108	10.456	11.382	7.001	17.058	10. 307	7.980	11.111
15						5.876	13.456	12.076	5. 825
	HEMBA1001225	13. 961	14.077	13.384	5. 925				
	HEMBA1001226	173.501	137.685	444.754	120.060	113.306	75. 167	63.960	67. 304
	HEMBA1001228	115.971	48.677	102.518	36.755	64.214	50.002	60.915	35. 178
	HEMBA1001229	246.802	111.161	135.886	43.460	94.703	148. 387	156.871	115.302
	HEMBA1001235	43.880	86.102	81.818	36.769	54. 172	65.830	70.065	66.201
20	HEMBA1001238	67.342	62.561	136.273	36.471	33. 852	41.838	26.195	28.747
20	HEMBA1001242	55. 562	43.106	58.593	41.382	47. 200	38.498	43.114	44.230
	HEMBA1001247	28.768	22.129	16.518	10.576	8.758	17.031	9, 651	13.385
	HEMBA1001253	58.130	80.415	66.640	18. 982	45. 992	54,071	95.073	63, 393
	HEMBA1001257			24. 256	10.657	12. 732	31.261	24.849	9, 134
		33.557	18.509						125.796
	HEMBA1001261	585. 214	143.415	243.791	98. 186	169. 988	310.109	234. 388	
25	HEMBA1001262	27.335	17.339	19.088	5. 647	15.678	20.899	11.464	19.889
25	HEMBA1001265	36.604	28.090	152.221	27.730	49.893	34. 423	16.502	26.993
	HEMBA1001266	69.367	67.414	170.657	45.898	31.802	39.554	41.287	52.480
	HEMBA1001269	69.921	44.649	36.964	34. 126	22. 232	42. 207	49.848	39.719
	HEMBA1001272	20.406	15.416	11.514	7.843	8.604	7.893	20.960	13.545
	HEMBA1001279	113.597	76.085	147.371	41.113	50.841	58. 248	43.344	47.548
	HEMBA1001281	45. 326	37.551	65. 225	44. 536	46.787	41.371	32.229	56.625
30	HEEBA1001286	370.697	150.949	236.623	103.571	123.976	219. 461	196.233	117.566
								26.637	19.344
	HEMBA1001289	41.041	24.670	40.151	15. 175	30.612	27.627		
	HEMBA1001291	76.537	40.444	50.226	18.776	38. 423	55. 355	46.692	35.972
	HEMBA1001294	82.258	72.319	157.642	42.143	20.735	29, 333	17.711	34.443
	HEMBA1001296	53.487	17.150	31.045	10. 275	15. 918	21.120	15.842	13.595
	HEMBA1001297	13.397	24.306	19.513	11.631	14.701	4. 543	9.800	8.121
35	HEMBA1001259	122.378	135, 140	326.747	90.817	73.749	56.152	49.803	80.999
	HEMBA1001302	56.839	29.036	56.412	19.108	20.078	34. 481	51.929	37.087
	HEMBA1001303	14. 975	18.442	43.778	16.797	10.985	11.442	9.787	19. 264
	HEMBA1001306	262.869	135.864	244. 234	109. 949	109, 582	147. 334	146.509	115.543
								56, 171	58.657
	HEMBA 1001308	174.017	96.705	220.049	56.953	61. 486 38. 245	74. 225	49. 204	51.006
	HEMBA1001310	103.029	52.915	67.714	22.895		67.233		
.40	HEMBA1001312	98.664	47. 333	61.080	18.118	33.555	47.007	41.795	38. 627
•	HEMBA1001319	2.396	8.234	13.960	1.828	5. 485	3.003	5.682	3.780
	HELBA1001322	139.794	39.912	105.709	27.700	41.977	70. 428	70.602	46. 470
	HEMBA1001323	33.347	16.728	25.356	11.399	17.982	11.181	6.356	12.033
	HEMBA1001326	86.190	37, 984	69.933	24. 331	30, 078	49.223	46.365	16.347
	HEMBA1001327	7. 232	9.387	23.180	7.314	5. 185	9, 563	4.423	5. 267
46	HEMBA1001330	115.768	106.951	275.315	73. 389	24, 661	70.535	40.088	77.680
45	HEMBA1001348				9. 575	13.666	23.703	12.647	13.724
		15.770	21.874	26. 347				34, 055	24.753
	HEMBA1001350	75.857	38.749	51.454	16. 428	34. 291	56.400		
	HEMBA1001351	52. 274	55.313	56.544	30. 521	46.408	29, 604	44.212	30.972
	HEMBA1001352	68.321	45.617	54.427	17. 559	29.887	39.484	52.789	
	HEMBA1001353	39.891	57.492	54.971	31. 425	27.945	45.687	29.741	66.188
50	HEPBA1001358	45, 659	52.406	59.774	46.865	40. 225	47.618	32.581	59.101
50	HEMBA1001361	22, 908	16.519	28.635	11.897	15, 569	13.635	13.938	
	HEMBA1001364		17. 205	23. 355	7.224	9.469	13.379	76. 125	
		18.896				25.778	40.077	21.715	
	HEMBA1001375	61.506	22.179	38.795	12.798				
	HEMBA1001377	140.430	131.029	307.084	83. 191	100.026	74.475	63. 988	
	HEMBA1001383	23. 974	26.206	28.704	11.442	17,819	19.160	16.899	
55	HELBA1001387	58. 343	34.130	63.677	19.556	30.371	42.397	40.247	49. 239

Table 18

	C				10 050 T	10 (24 )	20 022 1	22.224	33.218
	HEMBA1001388	48.601	24.690	39.877	18. 958	10,634	32.922		
	HEMBA1001390	132.003	94. 390	254.352	56.412	64.490	47, 169	44. 169	57.372
-	HEMBA1001391	18. 302	9, 686	12, 994	6.299	10.600	8. 500	7.116	5. 544
5	HEM8A1001398	91, 232	50. 992	142.408	36, 081	29. 548	29.490	28.704	29.984
	HEL:8A1001405	58.645	22.354	32. 227	15.864	9. 285	19, 993	24. 564	13.964
		36.434		105. 808	18.094	19. 994	13.316	18, 019	15.592
	HEMBA1001406		22.693			13. 107	23.014	18. 826	15.060
	HEMBA1001407	38.781	19.637	24. 599	18. 935				
	HEMBA1001411	28.412	7.180	21.950	8.303	9.708	14. 302	8. 598	6.663
10	HEMBA1001413	66.736	26. 480	35. 635	15.400	24.013	18.356	24. 304	20.769
,,	HEMBA1001414	20.720	7.567	18.414	12.522	9.722	12.903	18. 283	18.581
	HEMBA1001415	76.802	54, 702	159, 510	34, 156	20.989	32.235	21.694	26.676
	HEMBA1001416	41.784	23.474	29, 453	12.230	24.881	24. 993	25.847	28.651
	HEMBA1001432	74.066		190.870	40.409	63.619	36.879	66.751	33.675
	HEMBA1001433	132.672	110, 163	246. 542	77.852	61.676	50.447	37.821	64.403
		138.669	108.645	334. 104	89. 523	68.855	59.723	58.393	56.483
15	HEMBA1001435			11, 177	7. 985	15.704	7. 291	6.742	6.336
	HEMBA1001442	13.093	8, 604		40.086	27.976	37. 353	30. 266	41.647
	HEMBA1001446	102.450	63. 255	146.442			42.457	42.764	41.349
	HEMBA 1001450	72.339	35. 494	55. 103	30.799	31.322		41, 131	31.250
	HEMBA1001454	146.726	128.060	438. 247	88.679	43. 129	54.712		
	HEMBA 1001455	5.879	8. 197	8. 325	5. 561	4. 437	5. 252	4. 300	7. 359
20	HEMBA1001459	17. 432	15. 927	16. 490	6.749	2.733	5.888	7.836	10.963
~~	HEMBA1001461	61.531	52.734	57. 136	38.874	24.764	19.473	23. 241	32.318
	HEMBA1001462	10.875	14. 911	16, 843	12.984	13.465	48. 381	7.061	25. 992
	HEMBA1001463	137. 907	83.753	340.496	93.114	51.866	61.784	37.705	68.960
	HEMBA1001469	85.416	21,757	29. 463	15.911	84.887	77.440	27.033	29.537
	HEMBA1001473	20. 582	31.855	36.498	8.307	3.680	16.703	21.371	19.890
	HEMBA1001476	135.720	113.851	246, 800	65. 595	57, 431	63.903	65. 229	67.697
25	HEMBA1001477	5. 228	2,001	4. 505	2.645	1,540	3. 243	1.426	2.876
	HEMBA1001478	14. 335	10. 180	12.692	5. 468	4. 474	5. 444	2.171	4. 539
		88.891	28. 381	49. 689	21.660	14, 126	35.334	38.272	30, 563
	HEMBA1001480		5. 156	20. 900	4. 647	5, 264	9. 545	13.805	4, 424
	HEMBA1001483	29.872			7. 289	1. 585	5.016	5.792	5. 999
	HEMBA1001490	6.867	6. 967	14. 148			195. 947	194, 164	146. 945
30	KEMBA1001495	431.282	118.073	203.714	73. 985	176.836		23.206	45. 223
50	HE138A1001497	93.817	60. 807	227.867	55. 576	41.006	34. 182		76. 899
	HEMBA1001510	174. 254	120.414	343. 336	76.008	76. 932	73. 234	61.531	
	HEMBA1001515	45. 158	26. 337	67.169	15.756	15. 962	10.664	9.567	12.346
	HEMBA1001517	51.005	47.728	80. 287	34. 595	28. 246	21.020	17.229	33.972
	HEMBA1001522	7.431	8. 980	7.032	7.566	5.011	6.466	6.447	4. 824
	HEMBA1001526	48.774	21.300	32.732	18.831	22.395	22.767	23.530	17.914
35	HEMBA1001533	129, 423	85. 570	262.800	70.163	46.649	44.926	26.457	37. 421
	HEMBA1001547	59, 442	26, 656	27. 947	8.053	15. 558	53.508	108.861	25. 371
	HEPBA1001552	41.663	33, 242	115. 535	26. 222	30, 447	18.258	21.358	25.853
	HEMBA1001553	58, 388	75, 765	66.228	32.264	36. 396	54.513	64.874	41. 905
	HEMBA1001557	182, 516	80.827	161.852	69.344	80.644	123.765	111.732	70.946
	HEMBA1001563	39, 649	31, 429	85. 246	26.057	12, 157	15. 987	10.055	17:083
40	HELBA1001566	37.835	49.964	108. 284	35.793	23.255	25. 180	21.368	39.375
	HEMBA1001569	75. 584	44, 631	109.624	35. 487	130, 340	63, 130	44. 960	55. 257
	HEMBA1001570	198. 300	125. 319	444. 153	119.332	74. 267	79.979	64.732	90.896
			60.654	48.704	22.469	22.629	67.058	24.391	34. 300
	HEMBA1001579	103.128		312.570	131.687	142. 104	91.884	67.267	94.418
	HEMBA1001581	153.698	126. 225				4. 888	4. 671	5.144
.2	HEMBA1001582	3, 551	7.087	15. 302	4.019	8. 190	14. 993	7.628	12.297
45	HEM8A1001585	27. 271	18.375	25. 179	14. 108	5. 648			21.314
	HEMBA 1001589	109.877	22.722	49, 216	20. 427	22.904	64.665	57. 120	35. 562
	HEMBA1001595	71.600	62.349	46. 938	34. 447	29, 362	34.516	45. 233	
	HEYBA1001604	41.253	27.004	34. 167	16.004	6.061	21.932	18.414	23. 101
	HEMBA1001608	35.073	29. 270	41. 525	21.276	22. 867	22.699	14.094	15.366
	HEMBA1001615	556, 575	105.703	103.519	47.686	27.311	81.914	42.373	58.652
50	HEMBA1001620	134, 940	29.972	79.824	31.924	62.056	54. 423	64.359	35. 203
	HEMBA1001621	70.036	30.704	63, 807	15.048	19.545	42. 391	33.266	40.516
	HEE3A1001635	39. 932	29. 397	35.653	16. 214	18.765	19.655	22.405	14.095
	HELBA1001636	73.726	18. 596	35. 798	14. 928	12.865	24. 352	31.819	22.414
				79. 588	28. 452	22.449	25. 101	30.009	43.819
	HEMBA1001640	48. 402	45. 105		35. 084	26. 220	48.859	71.158	46.463
5.5	HEMBA1001647	82.402	39. 456	75. 907			208. 201	178. 161	96.640
55	HEMBA1001651	390.307	66.648	181.929	51.802	112.530	1 200. 201	1110.101	1 30.040

Table 19

	THE MOATAGE CEP	20 000		70 100 1	20 101	20 070	07 160	20 200 T	16 672
	HEMBA1001655	60.366	18. 983	58.438	20.404	25.072	27. 162	29. 260	26.673
	HEMBA1001658	6.754	15. 270	17.542	13.420	5.060	4.800	4.973	4.979
	HEMBA1001661	87.199	20.304	32.793	13.066	8.394	24.098	22, 916	24.583
5							86. 024		21.248
	HEMBA1001865	160. 583	20. 830	54.460	12.363	48.457		73.847	
	HEMBA1001670	16.953	38.651	17.002	34.999	14.855	17.849	22.906	29.478
	HEMBA1001672	32.013	18.885	29.000	10.798	7.763	13.782	17.314	12. 393
	HEMBA1001673		67, 401			14, 401	17.612	30.520	43, 461
		38. 188		34. 336	38.037				
	HEMBA1001675	25.652	15. 594	33.810	5.390	15.796	13. 173	20.020	12.830
40	HEMBA1001676	91.000	54.310	85.397	92.681	131.468	50. 365	47. 230	68.405
10	HEMBA1001678	218. 382	128. 995	336.408	93.889	115.305	80.843	48.879	83. 933
	TICHORIOTOIO						33. 368		
	HEMBA1001680	82. 159	51. 521	165.818	33.978	36.449		38. 495	35. 261
	HEMBA1001581	1.654	0.785	0.840	2.142	2.581	2.772	2.146	2.424
	HEMBA1001684	143. 985	84. 151	377.154	72.850	69.097	61.638	30.820	52.077
	HEMBA1001695	16.068	10.112	14.571	6.860	4.930	4.572	6.164	7. 330
								3.870	3,613
15	HEMBA1001702	26.509	13.637	8.186	8.466	4.041	2.043		
	HEMBA1001709	67.279	26. 552	35.845	13.982	21.742	28.610	24. 540	19.603
	HEMBA1001711	20.072	29. 559	39.037	20.902	21.639	12,713	14.718	33.127
	HEMBA1001712	80.448	25. 222	51.628	19, 393	12.482	38.014	39.474	14.831
						51.048		124.654	59. 543
	HEMBA1001714	360.368	55. 902	142. 225	33.748		144. 094		
	HEMBA1001717	78.599	137. 380	18.549	12.298	5. 575	38.689	10.120	6.047
00	HEMBA1001718	51.621	52. 280	151.597	31.305	21.166	29.146	14.075	24.411
20	HEMBA1001723	17.072	13.658	8.525	5.653	8.811	9. 350	11.097	7.268
	HELBA1001731		22.781		15, 151		15. 292	14. 020	16.584
		35.728		41.531		12. 421			
	HEMBA1001734	52. 546	40.599	99.556	25.099	24. 031	28. 537	17. 389	32.936
	HEMBA1001736	177. 269	58.328	110.046	33.820	58.955	108.630	91.464	62.571
	HEMBA1001741	41.432	12.649	29.883	14.886	16. 207	10.446	11.420	7.286
	HEMBA1001744			12.961	13, 191	14, 151	4.519	8.367	8.623
25		5, 531	6.849						
	HEMBA1001745	41.752	17.786	36. 239	12.476	21.118	23. 635	15.410	16.514
	HEMBA1001746	27. 437	14.874	24.099	8.668	21.929	19.488	11.306	10.070
	HEMBA1001761	93. 148	46.911	179.597	28.212	33.421	34.026	19. 164	25. 901
	HEMBA1001762	55. 612	45.069	102, 148	38.307	35.260	33. 316	21.274	45. 248
	HEMBA1001781	13. 298	21. 385	26, 693	6.898	17.098	52.601	11.768	23.068
	HEC3A1001784	89.965	43.765	70.054	26.575	31.708	50. 347	52. 265	31.618
30	HEMBA1001791	182. 379	81.719	171.066	44.628	49.350	82.856	58, 215	48. 207
	HEMBA1001794	248. 582	163.789	153,778	73.632	50.595	152. 279	178.827	132.329
									22.729
	HEMBA 1001800	23. 432	21.165	27.668	11.281	20.728	24. 910	36.900	
	HEMBA1001803	17.343	8. 333	22.801	6.620	6.043	7. 560	6.613	10.079
	HEMBA1001804	109, 775	44.797	59. 456	29.337	34.849	44.372	36.696	35.851
	HEMBA1001808	78. 129	23.567	38.056	15.858	23.507	27.136	14. 673	12.332
35							35.618	41.552	46, 141
••	HEMBA1001809	66. 887	31.733	54. 127	33, 314	26.179			
	HEMBA 1001811	58. 974	24. 196	37. 583	17.314	16.018	21.582	15.074	19.831
	HEMBA1001815	71.286	63,775	155, 707	37.153	29.944	35.297	25. 257	24. 172
	HEMBA1001816	38.494	19.017	16.797	7.139	5. 598	16.061	22. 304	14.646
	HEMBA1001819	18. 590	21.371	38. 109	20.938	21.358	15.313	14. 917	25.144
	HEMBA1001820	10.884	9. 530	8.017	3.507	4.470	3.473	2. 999	3.099
40	HEMBA1001822	74. 239	95.719	91.314	62.121	28. 285	42.988	38. 222	47.532
	HEMBA1001824	155. 543	93.583	301.248	95.135	67.478	89.045	64. 562	61.114
	HEMBA1001835	23.616	7.706	25.753	5.777	19.660	19.809	12.020	10.462
	HEMBA1001844	149. 876	52,023	230.213	48.968	42.113	39.652	33. 559	40.495
									35.814
	HEE3A1001847	52.045	19.220	40.636	20.235	5. 196	35. 109	20. 186	
	HEMBA1001849	101.048	104.708	250. 547	53.025	28.022	40.644	33. 371	35. 250
45	HEMBA1001850	105. 331	27.032	39.813	15.808	31.525	42, 751	44. 306	18.213
,,,	HEMBA1001861					4. 454	2. 945	0.995	3. 121
		3.104	4.469	6.763	3.292				
	HEMBA 1001862	50. 279	145. 708	102.412	25.750	34.563	40.833	22. 588	71.713
	HEMBA1001864	24.313	31.572	50.378	32.237	24.991	21.182	21.031	28. 126
	HEMBA1001866	57.711	54, 190	146,615	31.714	19.527	26.041	22.874	21.249
	HEMBA1001869					45. 195	40.562	22.644	40.891
	[UCTON 100 1893	55. 280	99. 559	58.454	35.799				
	(100.00)		44. 336	77, 195	41.540	39.300	54. 584	34. 598	42.631
50	HEMBA1001871	75.011							1 44 661
50				30, 568	85.092	19.827	15.356	8. 554	21.861
50	HEMBA1001876	34. 287	31.955	30.568					
50	HEMBA1001876 HEMBA1001878	34. 287 17. 361	31.955 17.619	17. 545	15.644	5.481	11.657	14.965	18.117
50	HEMBA1001876 HEMBA1001878 HEMBA1001879	34. 287 17. 361 57. 004	31.955 17.619 22.429	17. 545 37. 128	15.644 16.562	5. 481 20. 200	11.657 35.414	14. 965 21. 946	18.117 17.114
50	HEMBA1001876 HEMBA1001878 HEMBA1001879 HEMBA1001884	34. 287 17. 361	31.955 17.619	17. 545	15.644 16.562 38.470	5. 481 20. 200 27. 460	11.657	14.965	18.117
50	HEMBA1001876 HEMBA1001878 HEMBA1001879 HEMBA1001884	34. 287 17. 361 57. 004 68. 009	31.955 17.619 22.429 84.640	17. 545 37. 128	15.644 16.562 38.470	5. 481 20. 200 27. 460	11.657 35.414	14. 965 21. 946	18.117 17.114
	HEMBA1001876 HEMBA1001878 HEMBA1001879 HEMBA1001884 HEMBA1001886	34. 287 17. 361 57. 004 68. 009 12. 711	31.955 17.619 22.429 84.640 12.605	17. 545 37. 128 41. 930 37. 824	15.644 16.562 38.470 31.827	5. 481 20. 200 27. 460 15. 893	11.657 35.414 36.604 14.038	14. 965 21. 946 25. 345 6. 697	18.117 17.114 26.320 38.737
55	HEMBA1001876 HEMBA1001878 HEMBA1001879 HEMBA1001884	34. 287 17. 361 57. 004 68. 009	31.955 17.619 22.429 84.640	17. 545 37. 128 41. 930	15.644 16.562 38.470	5. 481 20. 200 27. 460	11.657 35.414 36.604	14. 965 21. 946 25. 345	18.117 17.114 26.320

Table 20

				22 7 2 7		0.4 460 1	00 COC T	20 720
HELEBA1001890	42. 902	42.848	42.779	30.112	25. 432	24. 430	22.605	26.730
HEMBA1001896	66.448	24.720	44. 103	21.972	17, 708	30.703	19.628	23.571
								20.813
HEMBA1001899	36. 251	25. 553	24. 121	14, 701	12.301	21.838	17. 455	
HEMBA1001904	54, 904	256.020	233.857	243, 646	55. 587	234.548	188. 571	526.744
				11.244	8. 226	15. 367	15.894	13.300
HEMBA1001910	40. 309	10.865						
HEMBA1001911	35. 962	23. 128	26. 357	25, 151	11.860	24. 224	22.870	18.238
HEMBA1001912	59. 924	66.966	97.679	51, 180	45. 903	33. 336	33.019	40.551
HEMBA1001913	175. 368	39.664	67. 432	33.132	26.376	63. 459	70.607	52.824
HEMBA1001915	14.756	14.666	30. 224	8. 295	7.629	17.718	6.737	8. 522
					أحسن والمستحدث	8. 265	4. 852	4.772
HEMBA1001918	5. 018	8. 961	27. 591	7.538	11.032			
HEMBA1001921	4. 431	8. 444	18.195	11.252	12, 587	7.417	7.668	2.769
HEMBA1001931	3.948	0.000	4.664	1.422	3. 480	2. 935	1. 127	2.898
HEMBA1001939	94. 821	24.679	81.706	24. 209	16. 592	37. 223	29.835	13.058
HEMBA1001940	54. 512	33. 931	145. 138	26.273	27.653	18. 649	13. 136	19.614
				18.718	14.782	25. 435	26.410	16, 143
HEMBA1001942	38. 572	16.710	32.402					
HEMBA1001944	210.898	71.197	96.883	48. 156	38. 533	82. 132	92.097	74.740
HEMBA1001945	31,531	17.019	14. 533	10.175	3.037	17. 421	12.222	11.694
								4.044
HEMBA1001950	7. 103	7.424	9.611	3. 281	4.091	7.632	5.310	
HEMBA1001951	46.024	19. 234	101.026	19. 207	13.212	23.714	20.006	19.402
HEMBA1001958	44. 554	12.806	35. 277	17. 321	13, 181	22.652	28.735	20.948
HEMBA1001960	20. 513	7.802	16.888	8. 822	2. 948	8. 826	10.834	12.935
HEMBA1001962	4. 367	5. 104	4. 205	2.811	3, 031	4.870	2.364	2.994
				18.757	6.663	17.775	- 8.068	8.601
HEMBA1001964	35. 944	22. 281	62.761					
HEMBA1001967	47. 345	29. 504	42.717	13.526	22.051	33. 555	23.601	37. 521
HEMBA1001979	35, 138	6.478	16.732	12.797	5.919	13.447	10.355	9.155
					24.898	26. 381	17.514	28.891
HEMBA1001987	60.083	52. 275	190. 331	45.735				
HEMBA1001991	111.286	79.833	276. 566	56.455	50. 862	50.789	40. 252	54.919
HEMBA1002003	66.389	23, 989	53.710	17.039	17, 174	30.547	28. 422	24.474
						24. 541	24. 237	27.345
HEMBA 1002005	86.885	41.457	150. 127	33.935	15. 339			
HEMBA 1002008	32. 101	25. 375	86.511	18. 349	8.912	7.593	18.519	14. 967
HEMBA1002018	66, 105	22, 380	36.174	16.334	21, 482	27.922	34.098	27.804
HEMBA1002022	13. 986	8.018	13.490	0.000	2. 985	5.730	6.036	1.433
HEMBA 1002029	132.547	305. 823	115. 974	144.692	70.087	74.071	37.046	204.730
	1 17 077	1 10 227	1 14 524	E 006	2 466	5 207 1	E 258	6 824
HEN8A1002030	17.077	10. 337	14. 524	5. 906	8. 466	5. 897	6. 258	6.824
HEMBA1002035	17.077 48.658	10. 337	14. 524	5. 906 14. 325	7.176	14.446	14.084	13.506
HEMBA1002035	48. 658	12.959	10.324	14, 325	7.176	14.446	14.084	13.506
HEMBA1002035 HEMBA1002037	48. 658 16. 343	12. 959 34. 097	10.324 27.567	14. 325 14. 451	7.176 12.568	14. 446 15. 087	14. 084 13. 819	13.506 12.140
HEMBA 1002035 HEMBA 1002037 HEMBA 1002038	48. 658 16. 343 68. 477	12. 959 34. 097 31. 733	10.324 27.567 91.391	14. 325 14. 451 16. 935	7. 176 12. 568 8. 370	14. 446 15. 087 6. 020	14. 084 13. 819 17. 500	13.506 12.140 19.367
HEMBA1002035 HEMBA1002037	48. 658 16. 343	12. 959 34. 097	10.324 27.567	14. 325 14. 451	7.176 12.568	14. 446 15. 087	14. 084 13. 819 17. 500 4. 716	13.506 12.140
HEMBA 1002035 HEMBA 1002037 HEMBA 1002038 HEMBA 1002039	48. 658 16. 343 68. 477 15. 944	12. 959 34. 097 31. 733 22. 707	10. 324 27. 567 91. 391 17. 807	14. 325 14. 451 16. 935 13. 914	7.176 12.568 8.370 7.910	14. 446 15. 087 6. 020 3. 306	14. 084 13. 819 17. 500 4. 716	13.506 12.140 19.367 11.003
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042	48. 658 16. 343 68. 477 15. 944 41. 657	12. 959 34. 097 31. 733 22. 707 27. 877	10. 324 27. 567 91. 391 17. 807 32. 654	14. 325 14. 451 16. 935 13. 914 21. 111	7. 176 12. 568 8. 370 7. 910 14. 815	14. 446 15. 087 6. 020 3. 306 10. 217	14. 084 13. 819 17. 500 4. 716 24. 300	13.506 12.140 19.367 11.003 22.659
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002042	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242	13.506 12.140 19.367 11.003 22.659 61.829
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042	48. 658 16. 343 68. 477 15. 944 41. 657	12. 959 34. 097 31. 733 22. 707 27. 877	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279	14. 325 14. 451 16. 935 13. 914 21. 111	7. 176 12. 568 8. 370 7. 910 14. 815	14. 446 15. 087 6. 020 3. 306 10. 217	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002043	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605	14.446 15.087 6.020 3.306 10.217 84.089 66.594	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002048	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417	12.959 34.097 31.733 22.707 27.877 92.912 29.889 84.099	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247	14.084 13.819 17.500 4.716 24.300 81.242 55.483 39.557	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002043	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441	14.084 13.819 17.500 4.716 24.300 81.242 55.483 39.557 25.376	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002048	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417	12.959 34.097 31.733 22.707 27.877 92.912 29.889 84.099	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002048 HEMBA1002053 HEMBA1002055	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115	12, 959 34, 097 31, 733 22, 707 27, 877 92, 912 29, 889 84, 099 19, 194 34, 916	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297	14.084 13.819 17.500 4.716 24.300 81.242 55.483 39.557 25.376	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002055	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002044 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002055	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002055	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA10020561 HEMBA1002061	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002061 HEMBA1002060	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002060 HEMBA1002080	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 33.904 13.652	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002061 HEMBA1002060	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652 101.175 72.141	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002055 HEMBA1002055 HEMBA1002060 HEMBA1002060 HEMBA1002060 HEMBA1002084 HEMBA1002084	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652 101.175 72.141	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652 101.175 72.141	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770 20.451
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002044 HEMBA1002046 HEMBA1002045 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002084 HEMBA1002084 HEMBA1002084 HEMBA1002085 HEMBA1002085 HEMBA1002085 HEMBA1002085	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652 101.175 72.141 18.082	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770 20.451 67.352
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 39.904 13.652 101.175 72.141	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770 20.451
HEMBA1002037 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002049 HEMBA1002053 HEMBA1002056	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139
HEMBA1002037 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 41.079 32.565 50.114	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459
HEMBA1002037 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002049 HEMBA1002053 HEMBA1002056	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565 50. 114	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 33.904 13.652 101.175 72.141 18.082 89.713 38.238 53.399 29.842	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960
HEMBA1002035 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002065 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002092 HEMBA1002092 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565 50. 114	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 33.904 13.652 101.175 72.141 18.082 89.713 38.238 53.399 29.842	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459
HEMBA1002035 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002042 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002051 HEMBA1002055 HEMBA1002055 HEMBA1002051 HEMBA1002051 HEMBA1002061 HEMBA1002100 HEMBA1002100 HEMBA1002101 HEMBA1002101	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565 50. 114 14. 867 31. 808	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 33.904 13.652 101.175 72.141 18.082 89.713 38.238 53.399 29.842 52.057	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002090 HEMBA1002090 HEMBA1002090 HEMBA1002100 HEMBA1002100 HEMBA1002101 HEMBA1002101 HEMBA1002107 HEMBA1002107	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 15. 801 14. 131 28. 199 321. 385	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 25. 842 52. 057 391. 825	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013	13.506 12.140 19.367 11.003 22.659 61.829 30.137 53.676 27.152 39.067 12.538 8.124 53.581 5.555 14.011 27.770 20.451 67.352 74.139 54.459 23.960 77.437
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002038 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002048 HEMBA1002053 HEMBA1002055 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002090 HEMBA1002090 HEMBA1002090 HEMBA1002100 HEMBA1002100 HEMBA1002101 HEMBA1002101 HEMBA1002107 HEMBA1002107	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565 50. 114 14. 867 31. 808	14.446 15.087 6.020 3.306 10.217 84.089 66.594 48.247 16.441 25.297 8.564 11.179 33.904 13.652 101.175 72.141 18.082 89.713 38.238 53.399 29.842 52.057	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002048 HEMBA1002048 HEMBA1002049 HEMBA1002053 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002090 HEMBA1002090 HEMBA1002100 HEMBA1002100 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002103 HEMBA1002103 HEMBA1002103 HEMBA1002103 HEMBA1002103 HEMBA1002103 HEMBA1002103 HEMBA1002103	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 4.867 31.808 369.500	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 42. 894 163. 850 236. 013 26. 533	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611
HEMBA1002035 HEMBA1002037 HEMBA1002038 HEMBA1002039 HEMBA1002042 HEMBA1002044 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002056 HEMBA1002105 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002101	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 106	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 72. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 28. 199 321. 385 18. 292 15. 858	7. 176 12. 568 8. 370 7. 910 14. 815 53. 861 21. 605 87. 434 16. 358 17. 449 1. 311 10. 973 72. 416 10. 773 19. 083 32. 654 15. 632 41. 079 32. 565 50. 114 14. 867 31. 808 369. 500 19. 991 20. 474	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217	14.084 13.819 17.500 4.716 24.300 81.242 55.483 39.557 25.376 28.606 4.481 9.701 45.653 6.835 43.117 50.433 12.882 44.294 15.932 40.628 22.894 163.850 236.013 26.533 28.894	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563
HEMBA1002037 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106
HEMBA1002037 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 106 84. 269	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 985 72. 351 45. 440 46. 944	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563
HEMBA1002035 HEMBA1002037 HEMBA1002037 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002049 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002101 HEMBA1002103	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 106 84. 269 37. 736	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440 46. 944 27. 034	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807 45. 990	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381 21. 694	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691 20. 410	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106 24. 305
HEMBA1002035 HEMBA1002037 HEMBA1002037 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 105 84. 269 37. 736 25. 756	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512 19. 103	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440 46. 944 27. 034 20. 941	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807 45. 990 4. 978	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311 11.161	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381 21. 694 9. 451	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691 20. 410 7. 795	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106 24. 305 9. 431
HEMBA1002035 HEMBA1002037 HEMBA1002037 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 105 84. 269 37. 736 25. 756	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512 19. 103	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440 46. 944 27. 034	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807 45. 990	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381 21. 694	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691 20. 410	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106 24. 305 9. 431 10. 190
HEMBA1002035 HEMBA1002037 HEMBA1002039 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002056 HEMBA1002061 HEMBA1002061 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002080 HEMBA1002103	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 106 84. 269 37. 736 25. 756 20. 036	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512 19. 103 10. 925 14. 349	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440 46. 944 27. 034 20. 941 19. 713	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 44. 136 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807 45. 990 4. 978 9. 608	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311 11.161 11.839 3.638	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381 21. 594 9. 451	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691 20. 410 7. 795	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106 24. 305 9. 431 10. 190
HEMBA1002035 HEMBA1002037 HEMBA1002037 HEMBA1002038 HEMBA1002043 HEMBA1002043 HEMBA1002043 HEMBA1002048 HEMBA1002048 HEMBA1002055 HEMBA1002055 HEMBA1002056 HEMBA1002105	48. 658 16. 343 68. 477 15. 944 41. 657 149. 364 137. 253 98. 417 33. 636 67. 115 13. 684 11. 815 59. 350 11. 331 69. 868 127. 409 34. 645 118. 301 57. 160 104. 746 35. 380 62. 621 745. 018 35. 812 42. 105 84. 269 37. 736 25. 756	12. 959 34. 097 31. 733 22. 707 27. 877 92. 912 29. 889 84. 099 19. 194 34. 916 12. 039 14. 960 80. 319 7. 502 62. 174 33. 016 16. 695 90. 733 69. 427 76. 058 25. 812 45. 738 396. 517 23. 546 14. 033 29. 512 19. 103	10. 324 27. 567 91. 391 17. 807 32. 654 208. 642 60. 279 271. 170 25. 821 39. 511 16. 129 29. 478 81. 497 15. 981 111. 196 60. 924 25. 357 129. 453 106. 418 178. 766 31. 300 65. 486 1335. 986 72. 351 45. 440 46. 944 27. 034 20. 941	14. 325 14. 451 16. 935 13. 914 21. 111 70. 906 19. 894 63. 157 11. 890 37. 518 14. 136 10. 168 43. 371 7. 301 13. 760 24. 219 15. 741 60. 276 34. 067 45. 801 14. 131 28. 199 321. 385 18. 292 15. 858 12. 807 45. 990 4. 978	7.176 12.568 8.370 7.910 14.815 53.861 21.605 87.434 16.358 17.449 1.311 10.973 72.416 10.773 19.083 32.654 15.632 41.079 32.565 50.114 14.867 31.808 369.500 19.991 20.474 29.311 11.161	14. 446 15. 087 6. 020 3. 306 10. 217 84. 089 66. 594 48. 247 16. 441 25. 297 8. 564 11. 179 39. 904 13. 652 101. 175 72. 141 18. 082 89. 713 38. 238 53. 399 29. 842 52. 057 391. 825 18. 086 47. 217 40. 381 21. 694 9. 451	14. 084 13. 819 17. 500 4. 716 24. 300 81. 242 55. 483 39. 557 25. 376 28. 606 4. 481 9. 701 45. 653 6. 835 43. 117 50. 433 12. 882 44. 294 15. 932 40. 628 22. 894 163. 850 236. 013 26. 533 28. 894 49. 691 20. 410 7. 795	13. 506 12. 140 19. 367 11. 003 22. 659 61. 829 30. 137 53. 676 27. 152 39. 067 12. 538 8. 124 53. 581 5. 555 14. 011 27. 770 20. 451 67. 352 74. 139 54. 459 23. 960 77. 437 348. 025 25. 611 33. 563 37. 106 24. 305 9. 431

Table 21

HEBBATOZITA 115, 045 48, 848 97, 208 42, 412 85, 310 14, 25, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 27, 207 14, 207 1		Derina A A A A A A			22 222 1		46 310 T	67 667 1	01 212	45.988
HEBBA1002151   St. 410   19.140   11.868   10.122   7.9318   26.996   19.485   14.196     HEBBA10027151   118.225   31.167   44.382   21.446   21.743   47.426   40.520   16.558     HEBBA10027161   72.851   68.019   132.155   42.302   37.035   29.438   49.438   41.818     HEBBA10027161   72.851   68.019   132.155   42.302   37.035   29.438   49.438   41.818     HEBBA10027161   72.851   68.019   132.155   42.302   37.035   29.438   49.438   41.818     HEBBA10027162   122.516   68.019   132.155   42.302   37.035   29.438   49.438   41.818     HEBBA10027163   49.869   43.502   64.932   20.426   7.151   0.000   10.979   12.578     HEBBA10027164   10.221   59.044   77.464   68.539   19.476   19.464   41.618     HEBBA10027167   179.053   81.40   59.594   77.464   68.539   19.476   19.464   41.618     HEBBA10027173   177.522   47.918   197.006   36.833   45.164   29.478   20.169   23.078     HEBBA10027173   177.522   47.918   197.006   36.833   45.164   29.478   20.169   23.078     HEBBA10027179   100.895   25.141   44.156   25.857   77.903   28.133   22.634     HEBBA10027179   55.617   55.403   85.686   45.680   26.918   80.684   59.110   46.489     HEBBA10027189   55.403   85.686   45.680   26.918   80.684   59.110   46.489     HEBBA10027189   75.840   77.5840   44.5740   44.5750   44.5750   44.5750     HEBBA10027189   75.840   76.095   44.771   17.903   28.133   22.637   44.631     HEBBA10027189   75.840   76.095   44.771   17.903   28.133   27.637   44.631     HEBBA10027189   75.840   77.8540   44.5750   44.5750   44.630   44.638     HEBBA10027180   77.8540   44.5750   44		HEHBA1002147	135.045	48.848	87. 208	42.412	46. 318	67.257	83. 313	
REBIADOZISS   12,728   25,418   35,746   20,325   25,538   18,972   25,019   19,327   19,32		HEMBA1002150		89.434	182. 502	48.715				
HEBBA1002153   12,758   25,478   35,746   20,375   25,638   15,972   26,019   19,425     HEBBA1002150   16,654   114,853   33,241   90,651   77,047   63,857   41,633   55,418     HEBBA1002161   72,851   88,019   32,155   42,303   77,035   29,438   49,435   41,633   55,418     HEBBA1002162   127,516   82,989   307,464   88,589   51,141   55,742   37,823   54,952     HEBBA1002163   43,869   43,602   64,932   20,426   7,151   0,000   30,973   25,788     HEBBA1002164   10,271   55,044   77,408   32,652   19,476   33,400   43,639   25,409     HEBBA1002164   10,271   55,044   77,408   32,652   19,476   33,400   43,639   25,240     HEBBA1002167   139,053   18,430   59,693   11,037   30,991   50,002   48,620   13,488     HEBBA1002177   10,085   25,137   76,834   194,618   71,071   73,577   73,787   73,787   73,787     HEBBA1002177   10,085   25,141   41,676   25,857   17,903   28,153   22,867   44,993     HEBBA1002179   10,281   19,500   46,777   13,260   23,273   37,375   44,478   27,788     HEBBA1002179   10,281   19,500   46,777   13,260   23,273   37,375   44,478   27,788     HEBBA1002179   10,281   17,500   46,777   13,260   23,273   37,375   44,478   27,788     HEBBA1002179   10,281   17,500   46,777   13,260   23,273   37,375   44,478   27,789     HEBBA1002179   10,281   17,500   46,777   13,260   23,273   37,375   44,478   27,789     HEBBA1002179   10,281   17,500   46,777   13,260   47,798   27,798   47,789	5	HEMBA1002151	60, 410	19.140	11.868	10.122	7. 938	26. 996		14. 196
HEBBAIDOZISS   18, 226   31, 167   44, 392   21, 446   21, 741   47, 426   40, 620   16, 558   HEBBAIDOZISS   16, 625   31, 167   44, 392   21, 446   21, 741   47, 426   40, 620   16, 558   HEBBAIDOZISS   16, 625   981   312, 158   42, 392   37, 305   29, 438   49, 438   41, 818   41	J	HEMBA 1002 153		25 478	35, 746	20, 325	25, 638	15. 972	26.019	19.827
HEBSIAD02180   165, 554   114, 853   318, 241   90, 651   71, 047   63, 857   41, 533   55, 419   HEBSIAD02152   72, 516   86, 919   312, 155   42, 302   37, 035   29, 308   49, 436   1, 818   HEBSIAD02152   127, 516   62, 989   307, 464   68, 589   51, 141   55, 242   37, 823   54, 929   74, 921   74,										16.858
Helbis 1007161   77, 851										
HEBSITODY:182   72,7   516   62,7   58   307,464   68,589   51,141   55,742   37,827   34,927   54,947   54,9										
The										
		HEMBA1002162	122.516	62.989	307. 464	68.589				
HEBBA1002164   110, 221   59, 044   71, 406   32, 552   19, 476   39, 440   43, 533   52, 044     HEBBA1002167   113, 053   18, 430   69, 693   11, 057   30, 091   50, 202   48, 520   13, 648     HEBBA1002177   100, 395   25, 141   41, 576   28, 587   17, 903   28, 153   22, 687   41, 081     HEBBA1002178   102, 331   19, 500   46, 771   31, 290   32, 323   37, 856   44, 115   27, 390     HEBBA1002178   102, 331   19, 500   46, 771   31, 290   32, 232   37, 856   44, 115   27, 390     HEBBA1002178   58, 517   58, 603   85, 588   45, 580   26, 918   50, 684   59, 110   54, 489     HEBBA1002178   79, 413   28, 230   31, 285   43, 185   40, 481   527, 390     HEBBA1002178   79, 413   28, 230   31, 285   23, 275   21, 094   33, 295   36, 478   41, 52     HEBBA1002189   79, 413   28, 230   31, 285   23, 275   21, 094   33, 295   36, 478   41, 52     HEBBA1002191   149, 027   80, 755   149, 493   49, 599   42, 372   50, 095   35, 614   44, 488     HEBBA1002191   79, 78, 785   28, 585   28, 288   16, 254   22, 500   31, 843   32, 595   35, 614   44, 488     HEBBA1002195   57, 388   28, 515   52, 22, 98   62, 524   522   50, 00   31, 843   32, 595   52, 791     HEBBA1002195   79, 786   28, 515   52, 22, 98   62, 524   522   500   31, 843   32, 595   52, 791     HEBBA1002195   79, 786   28, 515   52, 22, 98   62, 524   62, 525   52, 526   52	10	HEMBA1002163	49.889	43.602	64. 932	20.426	7.151	0.000		
HERBAI002716   312.586   255.137   568.314   194.638   171.071   159.271   134.442   213.993     HERBAI002717   107.532   47.918   197.005   36.333   76.164   29.418   20.169   23.078     HERBAI002717   107.552   47.918   197.005   36.333   26.164   29.418   20.169   23.078     HERBAI002717   107.331   19.500   46.777   13.290   32.123   37.855   44.115   27.390     HERBAI002718   55.617   55.000   46.777   13.290   32.123   37.855   44.115   27.390     HERBAI002718   55.617   55.000   46.777   13.290   32.123   37.855   44.115   27.390     HERBAI002718   55.617   55.000   46.777   13.290   37.177   22.480   37.355     HERBAI002718   79.413   28.280   31.825   23.275   27.049   37.177   22.480   37.355     HERBAI002718   79.413   28.280   31.825   23.275   27.049   37.177   22.480   37.355     HERBAI002718   79.413   38.80   31.825   23.275   27.049   37.177   22.480   37.355     HERBAI002718   79.413   38.80   31.825   23.275   27.049   37.177   22.480   37.355     HERBAI002718   75.439   70.805   48.011   47.097   37.440   31.295   36.478   38.275     HERBAI002718   75.149   70.805   48.011   47.097   37.440   31.295   36.478   38.275     HERBAI002718   75.149   70.805   48.011   47.097   37.440   31.295   35.478   38.275     HERBAI002718   75.149   75.855   75.868   28.515   52.628   15.254   27.800   37.843   37.855   37.55   37.55     HERBAI002718   74.937   37.55   37.55   37.55   38.57   37.55   37	,,	HEMBA1002164	110, 221	59.044	71, 408	32,652	19, 476	39. 440	43.639	52.404
HEBAT1002178   119, 053							171.071	159, 271	134, 442	213, 993
REBIATIOR3779   137.552										
HEBBAT002178   100, 895   25, 141   41, 576   25, 857   17, 903   28, 153   22, 687   14, 081   HEBBAT002178   152, 331   19, 500   46, 717   13, 390   12, 232   37, 856, 44, 115   27, 190   HEBBAT002185   35, 617   56, 403   85, 686   45, 680   26, 918   60, 684   55, 110   64, 849   HEBBAT002188   73, 413   28, 280   31, 285   23, 775   21, 094   33, 295   36, 475   18, 236   HEBBAT002189   75, 449   77, 588   212, 644   43, 915   27, 204   32, 127   24, 680   37, 386   HEBBAT002181   75, 439   76, 807   76, 807   76, 807   77, 775   77, 780   77, 780   78, 785   77, 780   78, 785   77, 780   78, 785   78										
The										
HEBBA1002118   S.S. 617   S.S. 403   85.868   45.860   26.918   60.684   55.110   54.849   SEPBA1002118   75.438   71.858   71.858   71.858   72.844   43.915   72.049   32.172   24.80   32.386   HEBBA1002118   75.439   70.605   71.801   71.802   72.480   72.386   71.802										
HEBBA1002185	15	HEH8A1002178	102.831	19.500	46.717					
### PERSATOOZ 189		HEMBA1002179	55.617	56.403	85. 586	45.680	26.918	60.684	59, 110	
HERBATIODZIBS 79, 413 28, 280 31, 325 23, 275 21, 094 33, 295 36, 478 18, 236 HEBBATIODZIBS 53, 497 76, 509, 148, 011 47, 092 32, 460 30, 101 44, 751 30, 532 HECBBATIODZIBS 151, 252 24, 996 74, 221 11, 3793 42, 372 60, 095 35, 614 44, 448 HEBBATIODZIBS 151, 252 24, 996 74, 221 11, 337 16, 495 12, 778 5 075 13, 556 HEBBATIODZIBS 14, 884 12, 040 36, 633 16, 632 15, 648 12, 789 14, 289 14,		HEMBA 1002 185	85, 236	71, 958	212.844	43.915	27.049	32. 172	22.480	32.386
HEBBATIOD2193							21,094	33, 295	36.478	18.236
HEBAT002192 15: 127 24: 996 24: 821 15: 373 16: 495 12: 50: 095 35: 614 44: 348 HEBAT002195 37: 368 28: 535 52: 828 16: 254 22: 500 31: 843 32: 995 29: 791 HEBAT002195 37: 368 28: 535 52: 828 16: 254 22: 500 31: 843 32: 995 29: 791 HEBAT002195 41: 884 12: 040 36: 533 16: 532 15: 443 16: 808 12: 631 17: 451 HEBAT002195 41: 848 12: 040 36: 533 16: 532 15: 443 16: 808 12: 631 17: 451 HEBAT002204 9: 525 5: 141 14: 889 6: 784 4: 619 10: 508 27: 818 3: 410 HEBAT002204 9: 525 5: 141 14: 889 6: 784 4: 619 10: 508 27: 818 3: 410 HEBAT002204 8: 822 44: 10: 946 8: 937 74: 537 745 745 745 745 745 745 745 745 745 74										
HEBBA1002195 57.368 28.535 52.828 15.254 22.600 31.843 32.995 29.791 HEBBA1002199 4.837 13.539 27.878 15.748 17.426 10.539 17.451 HEBBA1002199 4.837 13.539 27.878 15.788 17.426 10.539 15.564 8.927 HEBBA1002204 9.525 5.141 14.869 6.784 4.519 10.508 27.818 9.410 HEBBA1002212 8.709 6.241 10.946 9.855 2.602 5.864 5.366 4.214 HEBBA1002217 50.814 62.759 64.668 59.460 28.990 37.379 29.953 64.813 HEBBA1002217 50.814 62.759 64.668 59.460 28.990 37.379 29.953 64.813 HEBBA1002217 50.814 62.759 64.668 59.460 28.990 37.379 29.953 64.813 HEBBA1002220 27.731 14.997 21.655 8.451 6.409 5.563 1.641 6.714 HEBBA1002217 50.814 62.759 64.668 59.460 28.990 37.379 29.953 64.813 HEBBA1002220 27.731 14.997 21.655 8.451 6.409 5.563 1.641 6.714 HEBBA1002221 75.989 79.527 79.169 45.309 54.852 28.856 14.142 10.1597 HEBBA1002227 75.058 17.589 418.739 11.259 12.259 13.599 15.279 19.577 19.169 45.309 54.852 28.856 14.142 10.1597 HEBBA1002237 47.252 49.329 124.721 12.838 24.807 23.399 15.399 26.185 HEBBA1002237 47.252 49.329 124.721 12.838 24.807 23.399 15.399 26.185 HEBBA1002237 47.252 49.329 124.721 12.838 24.807 23.399 15.399 26.185 HEBBA1002237 47.729 45.781 19.083 19.839 12.516 12.703 85.899 61.450 90.668 HEBBA1002239 48.436 19.578 38.228 12.877 19.838 12.956 12.709 29.556 14.142 10.1597 HEBBA1002237 47.729 45.781 19.838 14.242 30.449 12.238 18.331 14.286 HEBBA1002237 40.584 18.584 18.584 18.541 43.824 30.449 23.288 18.331 14.286 HEBBA1002239 48.436 19.578 38.228 12.857 12.838 12.257 71.19 4.455 3.711 HEBBA1002229 48.436 19.578 38.228 12.857 12.838 19.339 12.257 71.19 4.455 3.711 HEBBA1002229 48.436 19.578 38.228 12.877 30.097 53.759 26.09 27.737 HEBBA1002230 10.831 10.758 18.277 10.09 19.766 11.508 10.900 25.959 14.425 10.031 HEBBA1002230 10.831 10.758 18.277 19.576 11.608 10.900 25.959 18.639 11.424 10.559 14.825 10.031 HEBBA1002230 10.783 19.578 38.228 12.877 30.097 53.759 28.348 19.144 16.635 HEBBA10022317 30.733 34.891 17.227 19.776 11.608 10.900 25.959 10.876 37.379 19.776 11.608 10.900 25.959 10.876 37.379 12.288 18.391 14.286 HEBBA										
HEBBA1002195   57, 168   28, 835   52, 248   16, 254   22, 800   31, 843   12, 851   17, 451     HEBBA1002199   24, 937   13, 539   27, 878   15, 728   17, 426   10, 539   19, 664   8, 977     HEBBA1002208   80, 832   44, 154   68, 317   68, 994   37, 453   74, 064   81, 827   112, 820     HEBBA1002212   8, 709   6, 241   10, 946   9, 855   2, 602   5, 864   5, 366   4, 214     HEBBA1002217   50, 81, 21   21, 209   8, 31, 165   19, 157   20, 170   17, 045   19, 124   21, 605     HEBBA1002217   50, 81, 41   62, 759   64, 668   59, 460   28, 990   37, 379   29, 953   64, 481     HEBBA1002227   50, 81, 41   41, 49, 41, 41   41, 41, 41, 41, 41, 41, 41, 41, 41, 41,	20									
HERBATOD2219										
HEIBAT002204 9.525 5.41 14.869 6.784 4.619 10.508 27.818 9.410 HEIBAT0022108 80.832 44.154 68.317 68.994 37.453 74.064 81.827 112.820 HEIBAT002215 36.521 28.098 31.165 19.157 20.170 17.045 19.124 21.605 HEIBAT002215 36.521 28.098 31.165 19.157 20.170 17.045 19.124 21.605 HEIBAT002216 9.824 10.828 31.165 19.157 20.170 17.045 19.124 21.605 HEIBAT002220 27.731 14.997 21.655 8.451 6.409 5.663 1.641 6.714 HEIBAT002226 91.222 113.507 269.906 83.183 68.283 59.461 56.996 78.924 HEIBAT002227 97.518 117.589 418.739 112.916 121.703 83.889 63.450 90.658 HEIBAT002229 170.518 117.589 418.739 112.916 121.703 83.889 63.450 90.658 HEIBAT002221 77.79 45.281 81.81.41 43.824 30.449 54.328 62.401 55.767 HEIBAT002227 7.79 45.281 81.81.541 43.824 30.449 54.328 62.401 55.767 HEIBAT002257 6.344 5.787 15.404 4.338 1.225 7.119 4.456 3.711 HEIBAT002257 6.344 5.787 15.404 4.338 1.225 7.119 4.456 3.711 HEIBAT002265 55.937 30.786 32.787 15.704 15.80 13.851 18.391 14.286 HEIBAT002267 271 029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEIBAT002267 271 029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEIBAT002267 51.540 26.396 27.766 20.313 15.579 28.384 19.144 16.695 18.896 18.391 14.286 HEIBAT002267 271 029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEIBAT002267 271 029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEIBAT002267 51.540 26.396 27.766 20.313 15.579 28.384 19.144 16.695 18.896 18.900 25.599 14.425 10.031 18.898 18.391 14.286 18.896 18.890 25.599 14.425 10.031 18.898 18.391 14.286 18.896 18.890 25.599 14.44 56.893 18.893				12.040	36.633					
HEBBA1002218 80.832 44.154 68.317 68.994 37.453 74.064 81.827 112.820 HEBBA1002217 80.998 31.165 19.157 20.170 17.045 19.124 21.0055			24. 937	13.539	27.878					
		HEM8A1002204	9, 525	5, 141	14.869	5.784	4.619	10. 508	27.818	
HEPBA1002217					68, 317	68.994	37. 453	74.064	81.827	112.820
HEBBA1002215   36.521   28.098   31.165   19.157   20.170   17.045   19.124   21.605     HEBBA1002217   50.834   57.759   64.668   59.460   28.990   37.379   29.963   64.813     HEPBA1002226   91.222   113.507   269.906   85.183   68.283   59.461   56.996   78.924     HEBBA1002227   55.957   91.527   79.169   45.309   54.823   59.461   56.996   78.924     HEBBA1002227   170.518   117.589   418.739   112.516   121.703   85.889   63.450   90.668     HEBBA1002227   47.252   49.329   124.721   32.838   24.807   23.399   15.399   26.185     HEBBA1002227   103.363   107.010   190.330   54.740   72.381   50.451   45.873   70.581     HEBBA1002224   70.729   45.281   81.541   43.824   30.449   54.328   62.401   55.767     HEBBA1002223   63.346   107.010   190.330   54.740   72.381   10.451   45.873   70.581     HEBBA1002223   63.346   19.578   38.228   12.875   21.884   23.928   18.619   17.388     HEBBA1002225   48.436   19.578   38.228   12.875   21.884   23.928   18.619   17.388     HEBBA1002225   48.436   19.578   38.228   12.875   21.884   23.928   18.619   17.388     HEBBA1002226   48.897   17.077   19.776   11.508   10.900   25.959   14.425   10.031     HEBBA1002226   44.897   17.077   19.776   11.508   10.900   25.959   14.425   10.031     HEBBA1002200   45.449   29.229   34.095   19.879   8.778   25.461   22.168   3.907     HEBBA1002200   45.489   77.077   19.776   11.508   10.900   25.959   14.425   10.031     HEBBA1002301   15.283   48.105   92.138   43.064   48.204   66.899   80.872   58.027     HEBBA1002302   152.883   48.105   92.138   43.064   48.204   66.899   80.872   58.027     HEBBA1002301   10.743   9.992   10.155   43.095   19.879   8.778   25.461   22.168   31.907     HEBBA1002310   10.743   9.992   10.155   43.549   2.547   7.995   4.048   5.700     HEBBA1002313   10.868   29.538   63.800   15.812   36.228   50.321   45.600   28.278     HEBBA1002313   35.4195   35.458   31.890   32.148   11.559   35.490   29.410   33.449   21.452     HEBBA1002313   35.4195   35.458   31.890   32.188   31.890   3						9, 855	2, 602	5.864	5. 366	4.214
REMBA1002217   50.834   52.759   54.668   59.460   28.990   37.379   29.963   64.813     REPBA1002220   27.731   14.997   21.655   8.451   6.409   5.663   1.641   6.713     REPBA1002227   55.957   91.527   79.169   45.309   54.832   28.856   14.142   101.597     REBBA1002227   55.957   91.527   79.169   45.309   54.832   28.856   14.142   101.597     REBBA1002227   47.252   49.329   112.916   121.703   83.889   61.450   90.668     REBBA1002227   47.252   49.329   124.721   32.838   24.807   23.399   15.399   26.185     REBBA1002239   103.363   107.010   190.830   54.740   72.381   50.451   45.873   70.581     REBBA1002231   70.779   45.228   81.541   43.824   30.449   54.328   25.401   55.767     REBBA1002257   6.344   5.877   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002257   6.344   5.877   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002257   6.344   5.878   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002258   48.436   19.578   38.228   12.455   21.842   21.928   18.619   17.988     REBBA1002257   6.344   5.877   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002257   6.344   5.877   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002257   6.344   5.877   15.404   4.338   1.225   7.119   4.455   3.711     REBBA1002257   6.348   3.105   2.737   2.827   15.078   28.043   25.609   27.237     REBBA1002265   35.947   30.786   32.747   24.827   15.078   28.043   25.609   27.237     REBBA1002206   44.897   17.027   19.776   11.608   10.900   25.595   14.425   10.031     REBBA1002300   152.883   48.105   92.158   43.644   43.044   65.899   80.872   58.027     REBBA1002300   152.883   48.105   92.158   43.064   48.204   65.899   80.872   58.027     REBBA1002300   152.883   48.105   92.158   43.064   48.204   65.899   80.872   58.027     REBBA1002300   152.883   48.105   92.158   43.064   48.204   65.899   80.872   58.027     REBBA1002300   152.883   48.105   92.158   43.064   48.204   65.899   80.872   58.027     REBBA1002330   152.883   48.105   92.158   4	25								19, 124	21.605 -
HEPBA1002226 27, 731 14, 997 21, 655 8, 451 6, 409 5, 663 1, 641 6, 714 HEPBA1002227 55, 957 91, 527 79, 169 45, 309 54, 892 28, 856 14, 142 101, 597 HEPBA1002229 170, 518 117, 589 418, 739 112, 916 121, 703 85, 889 61, 450 90, 688 HEBBA1002237 47, 252 49, 329 124, 721 32, 838 24, 807 23, 399 15, 399 26, 185 HEPBA1002239 103, 363 107, 010 190, 830 54, 740 72, 381 50, 451 45, 873 70, 581 HEPBA1002239 103, 363 107, 010 190, 830 54, 740 72, 381 50, 451 45, 873 70, 581 HEPBA10022239 25, 589 27, 877 35, 744 16, 605 13, 851 18, 938 18, 391 14, 286 HEPBA1002237 6, 344 5, 787 15, 404 4, 318 1, 225 7, 119 4, 455 3, 711 HEPBA1002259 48, 436 19, 578 38, 228 12, 875 21, 884 23, 928 18, 619 17, 988 HEPBA10022259 271, 027 219, 564 645, 284 192, 491 147, 403 112, 552 83, 057 137, 280 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 24, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 405 18, 425 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022267 108, 413 102, 522 243, 566 58, 776 30, 997 53, 750 74, 099 29, 752 HEPBA10022302 152, 883 48, 105 92, 158 43, 064 48, 204 66, 899 80, 872 58, 027 HEPBA10022304 66, 409 29, 289 34, 095 19, 879 8, 778 26, 461 22, 366 13, 907 HEPBA10022307 100, 402 132, 737 29, 252 544, 612 24, 050 42, 355 39, 076 37, 573 HEPBA10022307 100, 402 132, 737 29, 252 54, 612 24, 050 42, 355 39, 076 37, 573 HEPBA10022307 107, 43 9, 992 10, 165 4, 549 92, 549 10, 434, 445 10,										
HEBBA1002228 91.222 113.507 269.906 85.183 68.283 59.461 56.996 78.924 HEBBA1002227 55.957 91.527 79.169 45.309 54.892 28.856 14.142 101.597 HEBBA1002237 170.518 117.589 418.739 112.916 121.703 85.889 61.455 90.668 HEBBA1002237 47.252 49.329 124.721 32.838 24.807 23.399 15.399 26.185 HEBBA1002239 103.363 107.010 190.830 54.740 72.381 50.451 45.873 70.581 HEBBA1002239 25.559 27.877 35.744 16.605 13.851 18.938 18.391 14.286 HEBBA1002253 25.559 27.877 35.744 16.605 13.851 18.938 18.391 14.286 HEBBA1002259 48.436 19.578 38.228 12.375 21.884 23.328 18.619 17.988 HEBBA1002257 63.344 5.787 15.404 4.338 1.225 7.119 4.455 3.711 HEBBA1002257 63.443 57.87 15.404 4.338 1.225 7.119 4.456 3.711 HEBBA1002257 63.443 57.87 15.404 4.338 1.225 7.119 4.456 3.711 HEBBA1002257 68.413 107.522 243.566 58.776 30.097 53.750 24.099 27.237 HEBBA1002265 56.947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEBBA1002267 108.413 107.522 243.566 58.776 30.097 53.750 24.099 29.752 HEBBA1002267 618.413 107.522 243.566 58.776 30.097 53.750 24.099 29.752 HEBBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEBBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEBBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.595 HEBBA1002200 46.449 29.289 34.095 19.879 8.778 26.461 22.368 13.907 HEBBA1002200 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002200 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002210 10.783 7.996 12.646 4.775 10.008 4.330 5.128 3.630 HEBBA1002230 10.783 7.996 12.646 4.775 10.008 4.330 5.128 3.630 HEBBA1002231 504.772 93.600 191.534 46.814 134.386 238.599 265.167 88.087 HEBBA1002313 53.492 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEBBA1002313 53.492 21.508 32.148 11.559 15.490 29.410 33.449 21.452 HEBBA1002333 354.195 154.556 211.807 41.753 10.008 4.330 5.128 3.630 HEBBA1002337 63.852 21.508 21.859 4.556 4.011 5.799 4.476 4.666 HEBBA1002337 64.882 27.886 21.899 4.056 4.011 5.799 4.476 4.666 HEBBA1002337 64.885 25.508 21.859 4.056 4.011 5.799 4.076										
REBBA 002227   55.557   91.527   79.169   45.309   54.892   28.856   14.142   101.597										
REBBA1002229   170. 518   117. 589   418. 739   112. 916   121. 703   85. 889   63. 450   90. 668     REBBA1002237   47. 252   49. 329   124. 721   32. 838   24. 807   23. 399   15. 399   26. 185     REBBA1002231   103. 363   107. 010   190. 830   54. 740   72. 381   50. 451   45. 873   70. 581     REBBA1002231   70. 729   45. 281   81. 541   43. 824   30. 449   54. 328   62. 401   55. 767     REBBA1002237   5. 559   27. 877   35. 744   16. 605   13. 851   13. 938   18. 391   14. 286     REBBA1002259   48. 436   19. 578   38. 228   12. 875   21. 884   23. 928   18. 619   17. 988     REBBA1002259   48. 436   19. 578   38. 228   12. 875   21. 884   23. 928   18. 619   17. 988     REBBA1002259   48. 436   19. 578   38. 228   12. 875   21. 884   23. 928   18. 619   17. 988     REBBA1002259   56. 947   30. 786   32. 747   24. 827   15. 078   28. 043   29. 609   27. 237     REBBA1002265   56. 947   30. 786   32. 747   24. 827   15. 078   28. 043   29. 609   27. 237     REBBA1002267   108. 413   102. 522   243. 556   58. 776   30. 097   53. 750   24. 099   29. 752     REBBA1002260   44. 837   17. 027   19. 776   11. 608   10. 900   25. 959   14. 425   10. 031     REBBA1002302   152. 883   48. 105   92. 158   43. 064   48. 204   66. 899   80. 877   53. 502     REBBA1002302   152. 883   48. 105   92. 158   43. 064   48. 204   66. 899   80. 877   53. 502     REBBA1002302   152. 883   48. 105   92. 158   43. 064   48. 204   66. 899   80. 877   55. 027     REBBA1002302   10. 743   9. 992   10. 165   45. 81   14. 336   238. 599   56. 167   88. 087     REBBA1002319   2. 868   2. 455   9. 670   0. 933   4. 715   4. 369   5. 615   4. 579     REBBA1002328   89. 382   28. 578   41. 755   17. 175   20. 280   46. 772   34. 722   18. 301     REBBA1002333   54. 195   154. 586   21. 807   41. 794   124. 733   173. 522   284. 831   192. 502     REBBA1002334   6. 882   4. 859   8. 859   8. 859   51. 840   29. 410   33. 449   21. 452     REBBA1002335   14. 497   13. 001   12. 249   10. 426   11. 840   13. 977   71. 141   15. 760     R										
HEBBA1002237										
HEBRA1002239		HEMBA1002229								
HEMBAT002231 70.729 45.281 81.541 43.824 30.449 54.328 62.401 55.767 HEBBAT002232 25.559 27.877 35.744 16.605 13.851 83.938 18.391 14.286 3.711 HEMBAT002255 6.344 5.787 15.404 4.338 1.225 7.119 4.456 3.711 HEMBAT002259 48.436 19.578 38.228 12.875 21.884 23.928 18.619 17.988 HEMBAT002262 271.029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEMBAT002265 56.947 30.786 32.747 128.27 15.078 28.043 29.609 27.237 HEMBAT002265 56.947 30.786 32.747 128.27 15.078 28.043 29.609 27.237 HEMBAT002265 56.947 30.786 32.747 128.27 15.078 28.043 29.609 27.237 HEMBAT002265 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEMBAT002205 46.489 77.027 19.776 11.608 10.900 25.959 14.425 10.031 HEMBAT002205 15.288 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBAT002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 14.8105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBATO02302 10.402 132.737 29.225 24.012 24.050 42.355 39.076 37.573 HEMBATO02316 504.772 93.620 191.534 46.814 134.386 238.599 255.167 88.087 HEMBATO02317 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 FEMBATO02317 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 FEMBATO02317 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 FEMBATO02318 89.382 28.578 41.751 17.175 20.280 46.772 34.048 5.700 FEMBATO02317 10.743 9.992 10.165 4.549 2.547 7.952 4.048 5.700 FEMBATO02317 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 FEMBATO02313 354.195 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEMBATO02313 354.195 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEMBATO02313 16.488 29.538 63.800 15.812 36.228 50.321 45.600 28.278 10.000 23.278 10.000 23.278 40.000 2	30	HEMBA1002237	47. 252	49. 329	124.721		24. 807			
HEMBA1002257 25.559 27.877 35.744 16.605 13.851 8.938 14.286 3.711 HEMBA1002257 6.344 5.787 15.404 4.338 1.225 7.119 4.456 3.711 HEMBA1002257 48.436 19.578 38.228 12.875 21.884 23.928 18.619 17.988 HEMBA1002262 271.029 219.564 645.284 12.875 21.884 23.928 18.619 17.988 HEMBA1002265 56.947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEMBA1002265 56.947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEMBA1002265 56.947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEMBA1002205 51.540 26.395 27.765 20.313 15.579 28.348 19.144 16.695 HEMBA1002206 44.897 17.027 19.766 120.313 15.579 28.348 19.144 16.695 HEMBA1002206 44.897 17.027 19.766 11.608 10.990 25.959 14.425 10.031 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEMBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEMBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEMBA1002312 10.743 9.992 10.165 4.549 2.547 7.952 4.048 5.700 HEMBA1002313 63.542 21.208 22.148 11.559 15.400 29.410 33.449 21.455 HEMBA1002313 33.54 21.208 22.148 11.559 15.400 29.410 33.449 21.455 HEMBA1002313 63.542 21.208 22.148 11.559 15.400 29.410 33.449 21.455 HEMBA1002314 116.488 29.538 80.800 15.812 36.228 50.321 45.600 28.278 14.856 12.800 29.410 33.449 21.455 14.800 29.214 116.400 28.278 40.95 25.000 29.810 33.449 21.455 14.800 29.214 16.600 28.278 10.000 2348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEMBA1002315 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEMBA1002315 104.283 29.278 40.945 24.892 20.681 42.242 4		HEMBA1002239	103, 363	107,010	190.830	54.740	72.381	50.451	45.873	
HEBBA1002253 25.559 27.877 35.744 16.605 13.851 18.938 18.391 14.286 HEBBA1002257 6.344 5.787 15.404 4.338 1.225 7.119 4.456 3.711 HEBBA1002259 48.436 19.578 38.228 12.875 21.884 23.928 18.619 17.938 HEBBA1002262 271.029 219.564 645.284 192.491 147.403 112.552 83.057 137.280 HEBBA1002265 56.947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEBBA1002267 108.413 102.522 243.566 58.776 30.097 53.750 24.099 29.752 HEBBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEBBA1002286 44.897 17.027 19.776 11.608 10.900 25.959 14.425 10.031 HEBBA1002202 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEBBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 5.615 4.579 HEBBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 5.615 4.579 HEBBA1002313 36.542 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEBBA1002333 36.354 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.552 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.552 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.455 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.552 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.552 HEBBA1002333 354.155 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEBBA1002337 64.885 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEBBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.666 HEBBA1002348 6.388 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEBBA1002343 14.648 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEBBA1002348 6.388 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEBBA1002343 14.488 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEBBA1002345 6104.283 29.578 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.855 251.508 219.532 215.420				45, 281	81, 541	43.824	30.449	54. 328	62.401	55.767
HEMBA1002257   6.344   5.787   15.404   4.338   1.225   7.119   4.456   3.711     HEMBA1002259   48.436   19.578   38.228   12.875   21.884   23.928   18.619   17.988     HEMBA1002262   271.029   219.564   645.284   192.491   147.403   112.552   83.057   137.280     HEMBA1002265   56.947   30.786   32.747   24.827   15.078   28.043   29.609   27.237     HEMBA1002267   108.413   102.522   243.566   58.776   30.097   53.750   24.099   29.752     HEMBA1002270   51.540   26.395   27.766   20.313   15.579   28.348   19.144   16.695     HEMBA1002286   44.897   17.027   19.776   11.508   10.900   25.959   14.425   10.031     HEMBA1002202   152.883   48.105   92.158   43.064   48.204   66.899   80.872   58.027     HEMBA1002302   152.883   48.105   92.158   43.064   48.204   66.899   80.872   58.027     HEMBA1002304   6.050   6.814   19.492   7.905   4.038   7.098   5.307   1.737     HEMBA1002316   504.772   93.620   191.534   46.814   134.385   238.599   265.167   88.087     HEMBA1002319   2.868   2.456   9.670   0.933   4.715   4.369   5.615   4.579     HEMBA1002321   10.743   9.992   10.165   4.549   2.547   7.952   4.048   5.700     HEMBA1002322   10.743   9.992   10.165   4.549   2.547   7.952   4.048   5.700     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.499   21.452     HEMBA1002331   93.059   61.863   189.067   60.545   4.3745   40.085   13.954   34.456     HEMBA1002331   116.488   29.538   63.800   15.812   36.228   50.321   45.600   28.278     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.499   21.452     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.479   21.452     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.499   21.452     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.499   21.452     HEMBA1002333   63.542   21.208   32.148   11.559   15.490   29.410   33.499   21.452     HEMBA1002335   64.855   251.508   21.893   4.056   4.011   5.790   4.476   4.606     HEMBA1002357   6								18.938	18.391	14.286
HEBBA1002255										3, 711
HEBBA1002262   271   029   219   564   645   284   192   491   147   403   112   552   83   057   137   280     HEBBA1002265   56   947   30   786   32   747   24   827   15   078   28   043   29   609   27   237     HEBBA1002267   108   413   102   522   243   566   58   776   30   097   53   750   24   099   29   275     HEBBA1002286   44   897   17   027   19   776   11   608   10   900   25   959   14   425   10   031     HEBBA1002290   46   449   29   289   34   095   19   879   8   778   26   461   22   368   13   907     HEBBA1002302   152   883   48   105   92   158   43   064   48   204   66   899   80   872   58   027     HEBBA1002304   6   050   6   814   19   492   7   905   4   038   7   098   5   307   1   737     HEBBA1002316   504   772   93   620   191   534   46   814   134   386   238   599   265   167   88   087     HEBBA1002317   10   743   9   992   10   165   4   775   10   008   4   330   5   128   3   630     HEBBA1002337   89   382   28   578   41   753   17   175   20   280   46   772   34   722   18   301     HEBBA1002337   93   059   61   863   83   80   80   80   872   54   60   80   80   80   80   80   80   80										
HEEBA1002265 56, 947 30.786 32.747 24.827 15.078 28.043 29.609 27.237 HEEBA1002267 108.413 102.522 243.566 58.776 30.097 53.750 24.099 29.752 HEEBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEEBA1002286 44.897 17.027 19.776 11.608 10.900 25.959 14.425 10.031 HEEBA1002280 44.897 17.027 19.776 11.608 10.900 25.959 14.425 10.031 HEEBA1002280 46.449 29.289 34.095 19.879 8.778 26.461 22.368 13.907 HEEBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEEBA1002304 6.050 6.814 19.492 7.905 4.038 7.098 5.307 1.737 HEEBA1002307 100.402 132.737 29.225 24.612 24.050 42.355 39.076 37.573 HEEBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEEBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 5.615 4.579 HEEBA10023128 89.382 28.578 41.753 17.175 10.008 4.330 5.128 3.630 HEEBA1002328 89.382 28.578 41.753 17.175 20.280 46.772 34.722 18.301 HEEBA1002333 63.542 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEEBA1002333 354.195 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002355 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEEBA1002356 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002356 17.449 51.764 63.3278 67.711 43.280 33.755 31.248 49.484	25									
HEEBA1002267 108.413 102.522 243.566 58.776 30.097 53.750 24.099 29.752 HEEBA1002270 51.540 26.396 27.766 20.313 15.579 28.348 19.144 16.695 HEEBA1002286 44.897 17.027 19.776 11.608 10.900 25.959 14.425 10.031 HEEBA1002290 46.449 29.289 34.095 19.879 8.778 26.461 22.368 13.907 HEEBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEEBA1002304 6.050 6.814 19.492 7.905 4.038 7.098 5.307 1.737 HEEBA1002304 6.050 6.814 19.492 7.905 4.038 7.098 5.307 1.737 HEEBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEEBA1002319 2.868 2.456 9.670 0.933 4.715 4.359 5.615 4.579 HEEBA1002320 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 HEEBA1002321 10.743 9.992 10.165 4.549 2.547 7.952 4.048 5.700 HEEBA1002333 63.542 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEEBA1002333 354.195 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 6.088 1.306 3.748 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 6.088 1.306 3.748 HEEBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 6.088 1.306 3.748 HEEBA1002345 6.318 7.600 13.603 5.490 2.590 6.088 1.306 3.748 HEEBA1002345 6.318 7.600 13.603 5.490 2.590 6.088 1.306 3.748 HEEBA1002345 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEEBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.33	33									
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HEBBA1002302 152.883 48.105 92.158 43.064 48.204 66.899 80.872 58.027 HEBBA1002304 6.050 6.814 19.492 7.905 4.038 7.098 5.307 1.737 HEBBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEBBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 265.167 88.087 HEBBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 5.615 4.579 HEBBA1002312 10.743 9.992 10.165 4.549 2.547 7.952 4.048 5.700 HEBBA1002312 89.382 28.578 41.753 17.175 20.280 46.772 34.722 18.301 HEBBA1002313 63.542 21.208 32.148 11.559 15.490 29.410 33.449 21.452 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.456 HEBBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.456 HEBBA1002338 66.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEBBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEBBA1002356 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 71.449 51.764 63.278 52.711 43.280 33.755 31.248 49.484					34.095	19.879	8.778	26.461	22. 368	
HEBBA1002316 504.772 93.620 191.534 46.814 134.386 238.599 265.167 88.087 HEBBA1002319 2.868 2.456 9.670 0.933 4.715 4.369 5.615 4.579 HEBBA1002320 10.783 7.936 12.646 4.775 10.008 4.330 5.128 3.630 HEBBA1002312 10.743 9.992 10.165 4.549 2.547 7.952 4.048 5.700 HEBBA1002312 89.382 28.578 41.753 17.175 20.280 46.772 34.722 18.301 HEBBA1002317 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.456 HEBBA1002333 63.544 95 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEBBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEBBA1002355 104.283 29.278 40.945 18.593 4.056 4.011 5.790 4.476 4.606 HEBBA1002357 64.853 29.278 40.945 21.809 20.581 42.242 45.108 28.190 HEBBA1002357 64.883 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.883 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.883 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.883 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.883 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002357 74.499 51.764 63.278 52.711 43.280 33.755 31.248 49.484	40							66.899	80.872	58.027
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HEMBA1002337 93.059 61.863 189.067 60.545 43.745 40.085 13.954 34.456 HEMBA1002339 354.195 154.586 211.807 141.794 124.733 173.522 284.831 192.502 HEEMBA1002341 116.488 29.538 63.800 15.812 36.228 50.321 45.600 28.278 HEMBA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEMBA1002349 6.318 7.600 13.603 5.490 2.590 6.088 1.306 3.748 HEMBA1002353 14.497 13.001 12.249 10.426 11.840 13.977 17.141 16.760 HEEMBA1002356 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEMBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEMBA1002365 71.449 51.764 63.278 52.711 43.280 33.755 31.248 49.484		HEMBA1002333	63. 542	21.208	32.148	11.559	15. 490	29.410	33.449	21.452
HEMBA1002339 354. 195 154. 586 211. 807 141. 794 124. 733 173. 522 284. 831 192. 502 HEMBA1002341 116. 488 29. 538 63. 800 15. 812 36. 228 50. 321 45. 600 28. 278 HEMBA1002348 6. 882 4. 859 18. 593 4. 056 4. 011 5. 790 4. 476 4. 606 HEMBA1002349 6. 318 7. 600 13. 603 5. 490 2. 590 6. 088 1. 306 3. 748 HEMBA1002353 14. 497 13. 001 12. 249 10. 426 11. 840 13. 977 17. 141 15. 760 HEMBA1002356 104. 283 29. 278 40. 945 24. 892 20. 681 42. 242 45. 108 28. 190 HEMBA1002357 64. 855 251. 508 219. 532 215. 420 68. 836 206. 728 136. 339 380. 371 HEMBA1002365 71. 449 51. 764 63. 278 45. 271 43. 280 33. 755 31. 248 49. 484			93.059		189.067	60.545	43. 745	40.085		34.456
HEEBA1002341 116, 488 29, 538 63, 800 15, 812 36, 228 50, 321 45, 600 28, 278 HEEBA1002348 6, 882 4, 859 18, 593 4, 056 4, 011 5, 790 4, 476 4, 606 HEEBA1002349 6, 318 7, 600 13, 603 5, 490 2, 590 6, 088 1, 306 3, 748 HEEBA1002353 14, 497 13, 001 12, 249 10, 426 11, 840 13, 977 17, 141 15, 760 HEEBA1002355 104, 283 29, 278 40, 945 24, 892 20, 681 42, 242 45, 108 28, 190 HEBBA1002357 64, 855 251, 508 219, 532 215, 420 68, 836 206, 728 136, 339 380, 371 HEEBA1002365 87, 281 64, 882 77, 475 30, 773 56, 108 61, 060 59, 371 56, 291 HEEBA1002365 71, 449 51, 764 63, 278 52, 711 43, 280 33, 755 31, 248 49, 484								1 173. 522	284.831	192.502
HEBSA1002348 6.882 4.859 18.593 4.056 4.011 5.790 4.476 4.606 HEPSA1002349 6.318 7.600 13.603 5.490 2.590 6.088 1.306 3.748 HEBBA1002353 14.497 13.001 12.249 10.426 11.840 13.977 17.141 16.760 HEBBA1002356 104.283 29.278 40.945 24.892 20.681 42.242 45.108 28.190 HEBBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEBBA1002365 71.449 51.764 63.278 52.711 43.280 33.755 31.248 49.484										28. 278
HEPA100235 6. 318 7. 600 13. 603 5. 490 2. 590 6. 088 1. 306 3. 748 HEBA1002353 14. 497 13. 001 12. 249 10. 426 11. 840 13. 977 17. 141 16. 760 HEBA1002356 104. 283 29. 278 40. 945 24. 892 20. 681 42. 242 45. 108 28. 190 HEBA1002357 64. 855 251. 508 219. 532 215. 420 68. 835 206. 728 136. 339 380. 371 HEPA1002360 87. 281 64. 882 77. 475 30. 773 56. 108 61. 060 59. 371 56. 291 HEPA1002363 71. 449 51. 764 63. 278 52. 711 43. 280 33. 755 31. 248 49. 484										
HELBA1002353 14. 497 13. 001 12. 249 10. 426 11. 840 13. 977 17. 141 16. 760 HELBA1002356 104. 283 29. 278 40. 945 24. 892 20. 681 42. 242 45. 108 28. 190 HELBA1002357 64. 855 251. 508 219. 532 215. 420 68. 836 206. 728 136. 339 380. 371 HELBA1002360 87. 281 64. 882 77. 475 30. 773 56. 108 61. 060 59. 371 56. 291 HELBA1002362 71. 449 51. 764 63. 278 52. 711 43. 280 33. 755 31. 248 49. 484	50									
HEEBA1002356 104. 283 29. 278 40. 945 24. 892 20. 681 42. 242 45. 108 28. 190 HEBBA1002357 64. 855 251. 508 219. 532 215. 420 68. 835 206. 728 136. 339 380. 371 HEEBA1002360 87. 281 64. 882 77. 475 30. 773 56. 108 61. 060 59. 371 56. 291 HEEBA1002363 71. 449 51. 764 63. 278 52. 711 43. 280 33. 755 31. 248 49. 484										
HEHBA1002357 64. 855 251. 508 219. 532 215. 420 68. 835 206. 728 136. 339 380. 371 HEHBA1002360 87. 281 64. 882 77. 475 30. 773 56. 108 61. 060 59. 371 56. 291 HEHBA1002363 71. 449 51. 764 63. 278 52. 711 43. 280 33. 755 31. 248 49. 484		HEMBA 1002353								
HEHBA1002357 64.855 251.508 219.532 215.420 68.836 206.728 136.339 380.371 HEHBA1002360 87.281 64.882 77.475 30.773 56.108 61.060 59.371 56.291 HEHBA1002363 71.449 51.764 63.278 52.711 43.280 33.755 31.248 49.484		HEEBA1002356		29. 278	40.945	24.892	20. 681			
HEMBA1002360 87. 281 64. 882 77. 475 30. 773 56. 108 61. 060 59. 371 56. 291				251.508	219.532	215, 420	68. 836	206.728		
UEF-BA1002363 71 449 51 764 63 278 52 711 43 280 33 755 31 248 49 484					_		56. 108	61.060	59.371	56.291
55									31.248	49.484
	55	I EFFUI AATAA	1 . 1. 773	1 01.104		1 33	1			

Table 22

							-0.000	0.365
HEMBA1002365	13.435	10.346	9.534	5. 175	9.470	4. 445	10.802	9. 325
HEMBA1002370	29.997	4. 107	11.054	4, 163	3.224	9.009	7.477	2.921
					10.204	18.275	29. 203	18.856
HEMBA1002374	91.498	18.475	11. 325	15.862				
HEMBA1002376	185.416	75. 425	127.578	52, 056	38.450	99. 590	90. 190	86.994
HEMBA1002377	81.350	41.908	63.893	37. 221	23.657	110.374	162. 166	50.770
								127.767
HEMBA1002380	189.521	137.466		137. 908	491.500	90.431	81.778	
HEMBA1002381	195.037	101.891	447. 953	125. 938	88.330	90, 756	70, 293	106.965
			42. 496	14.694	19.780	40, 126	24. 243	12.399
HELBA1002384	35. 247	22.319						
HEMBA1002389	44.796	8.467	36.790	11.793	9.362	18.736	15.497	20.728
HEMBA1002396	101.267	69.467	33. 025	16.553	26.429	25. 964	22. 294	23.666
				8.848	9.913	21, 219	18.569	22.818
HEMBA1002402	75.818	24. 148	28. 457					
HEMBA1002417	132.807	33.708	84. 436	22.910	38.826	58, 589	58.836	38.486
HEMBA1002419	75, 547	31.202	41.690	13.558	16.457	27. 281	19.705	13.013
			35, 559	17.034	13.878	23.652	14.721	24,637
HEMBA1002420	20.818	20.448						
HEMBA1002421	23.903	25. 285	59.023	7.957	14.189	24, 230	61.011	21.849
HEMBA1002423	12.762	11.755	25, 941	12, 938	14.177	14, 263	12.495	7.512
					25.667	42.797	41.513	31.249
HEMBA1002424	111,995	32. 293	46.657	24. 424				
HEMBA1002426	60.617	23.489	45.906	20. 305	25. 173	30.860	37.738	21.223
HELBA1002430	24, 143	3. 128	4. 900	1.517	4. 594	3, 316	8. 552	3.069
	59.808	37.476	93. 025	16.789	23.324	47.857	33.099	27.888
HEMBA1002439								
HEMBA1002441	77.869	99. 262	110.341	38.723	34. 562	65. 309	85.421	66.581
HEMBA1002454	58. 292	15, 281	38. 384	7.520	19.044	25. 972	22.845	22.015
HEL BA1002458	57, 329	46.103	101.242	30, 906	82.184	61,800	25.094	59.039
***************************************							14.678	14, 249
HELBA1002460	32.814	9. 205	25. 085	12.160	23.009	18.683		
HEMBA1002462	98.420	38. 135	55. 208	10.919	24.257	49.697	43.851	32.387
HEMBA1002465	11.819	15.260	28. 272	11.939	11.225	10.938	13. 593	20.635
				39. 999	39.213	76, 320	69.012	86.309
HEMBA1002469	129.538	61.348	120. 187					
HEMBA1002475	3.180	5.116	4. 323	2.230	1.467	4. 495	10.058	15.691
HEMBA1002477	93.696	64,730	238.114	55. 207	43, 349	42.487	29.532	52.786
HELBA1002480	210.023	58. 823	84. 566	37.478	45.060	106.554	97.791	70, 487
							42. 334	82.875
HEL'8A1002481	104.499	76.474	222. 903	71.502	68.097	67.421		
HEMBA1002486	81.465	42.269	169. 291	49.953	40.852	39.475	29. 153	26. 233
HEMBA1002490	66.695	11, 331	31.314	14, 602	25.852	35, 945	35. 954	15. 278
		12.315	25. 235	7.937	4.091	17.402	14. 269	10.773
HEMBA1002495	59. 387							12.219
HELTBA1002498	56.425	23.969	67.108	11.632	15.655	24.420	8. 272	
HEMBA1002501	40.955	16.994	22.074	1 13. 575	16.498	21.707	39.506	24. 519
HEMBA1002503	81.763	65.044	154. 595	39.638	33.778	31.214	32.219	26.800
							52.190	53. 323
HEMBA1002504	155. 357	95.219	279. 391	90.092	120.246	70.516		
HEMBA1002508	99.443	88. 234	259.961	107.085	79.039	59. 181	59.924	61.423
HEMBA1002513	50.560	22.902	30. 431	26.184	20.783	30,500	32.903	22.864
						20.264	20.643	13.727
HEMBA1002515	60.938	23.064	25. 098	16, 172	5.716			
HEMBA1002524	94.350	36.789	56.675	25.998	28.978	49.840	57.148	25. 205
HEMBA1002538	116,609	19.632	26.764	12.798	20.203	16.422	17. 588	15.759
HEPBA1002542	81.641	81.952	188.888	54. 986	41.864	32.890	30.719	38. 321
								21.614
HEMBA1002544	52.394	49. 175	98.415	47. 569	28. 375	28.766	20.948	
HEMBA1002546	76.538	62.763	156.051	47.625	74.374	45. 975	34.756	46.753
HEMBA1002547	11.448	4, 516	10.647	4.733	12.220	11.801	9, 959	7.127
HEYBA1002550	67. 373	39.322	48. 468	15.671	16.497	121.814	94, 586	25. 401
							16.656	18.665
HEL BA1002551	94, 391	14.109	27.085	11.976	8.787	41.811		
HEMBA1002552	204, 583	77.430	205. 444	49.448	44.756	67.408	63.216	57.684
HEMBA1002555	25, 583				5.608	14.795	10.111	7.416
				59.079	41.247	33. 253	41.617	41.270
HEMBA1002558	92.744	77.405						
HELBA1002561	53.810	51.725	155. 895	34.956	27.689	17.264	10.138	27. 24
MEMBA1002562	15. 261	10.822	15. 435	8. 259	18.723	12.036	9.056	10. 429
HEMBA1002568	24. 946	17.442	35, 354	17.552	10. 575	15, 262	16, 158	22. 328
						54, 936		27. 309
HEMBA1002559	112.340	34. 133	118.192	37.823	57.431		26. 164	
HEMBA1002570	43. 528	50.809	52.195	34.901	23.728	28. 874	9.812	50. 494
HEMBA1002574	106, 101	25. 148		16.369	26.322	57. 278	42.795	31.310
						1 20, 158	16.837	16.418
HEMBA 1002583	36.042	17.582		12.456	13.418	4		
HEMBA1002587	61.527	32.123	45.811	22.217	18.974	32.461	40. 250	39.915
HEMBA1002590	151.583	106.074	287.276	84.766	32.321	58. 221	38.642	53. 855
						53. 653	35. 724	57.578
HEMBA1002592	97.854			89. 335				
HEMBA 1002595	146.016	25.688		24. 156		86.770	79.174	26.760
HEMBA1002609	97.442	41. 925	56.054	29.427	35.650	35.839	44.588	47.074
11 INCHEST   VALUES	1 776		1 22.007		1			·

Table 23

HEBBAIDOZEST 1 75.792 85.517 99.445 13.277 12.399 21.055 16.512 49.186 HEBBAIDOZEST 1 01.131 25.998 30.99 15.921 1.913 0.018 4 35.003 28.108 HEBBAIDOZEZ 1 14.592 25.865 18.082 8.927 7.391 8.869 5.823 12.263 HEBBAIDOZEZ 2 194.515 18.082 8.927 7.391 8.869 5.823 12.263 HEBBAIDOZEZ 2 194.515 18.082 8.927 7.391 8.869 10.257 7.527 11.624 HEBBAIDOZEZ 3 12.199 16.170 19.306 15.884 8.0306 7.1673 11.673 11.3228 101.786 5.515 14.1673 11.3228 101.786 5.515 14.1673 11.3228 101.786 5.515 14.1673 11.3228 101.786 5.515 14.1673 11.3228 101.786 5.515 14.1673 11.3228 11.3228 101.786 5.515 14.1673 11.3228 11.32		•			iao	ne 23				
HEIBAIDOZES   14,592   25,365   80,999   15,921   21,913   40,814   15,003   28,108   16,003   16,003   16,003   17,003   18,108   18,003   17,003   18,108   18,003   17,003   18,108   18,003   17,003   18,108   18,003   17,003   11,022   10,1786   53,514   17,003   17,003   17,003   11,022   10,1786   53,514   17,003   17,003   17,003   11,022   10,1786   53,514   17,003   17		HEMRA 1002617	26.792	86 617 1	59, 446	73.277	12.909	21.055	16.612	49. 136
HEBBAIDOZEZ   14,582   25,865   18,082   8,977   7,391   8,869   5,823   12,283     HEBBAIDOZEZ   136,461   21,509   21,649   9,956   16,559   10,257   7,527   11,624     HEBBAIDOZEZ   12,199   13,700   29,060   15,884   27,720   10,277   7,527   11,624     HEBBAIDOZEZ   12,199   18,700   29,060   15,884   27,840   28,811   37,912   40,046     HEBBAIDOZEZ   15,526   48,044   90,956   15,894   27,840   28,811   37,912   40,046     HEBBAIDOZEZ   17,869   24,187   46,646   22,248   22,950   37,216   25,867     HEBBAIDOZEZ   17,869   24,187   46,646   22,248   22,950   37,216   25,867   21,387     HEBBAIDOZEZ   17,869   24,187   46,646   22,248   22,950   37,216   25,867   21,387     HEBBAIDOZEZ   17,869   24,187   46,646   22,248   22,950   37,216   25,827   21,387     HEBBAIDOZES   13,382   27,730   33,313   16,958   11,617   23,904   29,214   16,599     HEBBAIDOZES   13,382   27,730   33,313   16,958   11,617   23,904   29,214   16,599     HEBBAIDOZES   18,498   86,104   154,110   51,966   22,499   22,990   27,314   23,777     HEBBAIDOZES   15,384   86,104   154,110   51,966   22,499   22,990   27,314   23,777     HEBBAIDOZES   15,384   86,144   54,110   51,966   27,580   20,804   21,902   21,3470     HEBBAIDOZES   15,384   86,144   54,110   51,966   27,580   20,804   21,902   21,3470     HEBBAIDOZES   15,384   86,244   64,689   29,523   20,013   30,244   25,612   28,3769     HEBBAIDOZES   16,631   88,966   247,534   44,722   23,400   33,514   5,613   22,532     HEBBAIDOZES   16,631   88,966   247,533   44,723   44,725   23,400   44,725     HEBBAIDOZES   15,534   64,138   56,537   37,679   18,172   23,470   38,288   44,113     HEBBAIDOZES   15,534   64,138   36,653   37,343   31,475   34,472   34,472   34,472     HEBBAIDOZES   15,534   64,138   34,472   34,472   34,472   34,472   34,472     HEBBAIDOZES   15,534   64,535   67,56   73,679   18,172   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,470   34,						15.921	21.913	40.814	35.003	28. 108
HEBBAIDOZEZ   254, 635   47, 837   73, 568   48, 036   71, 673   713, 228   101, 786   53, 514     HEBBAIDOZEZ   31, 044   47, 279   72, 504   73, 506   16, 559   10, 227   7, 527   11, 624     HEBBAIDOZEZ   52, 796   46, 044   90, 968   58, 904   72, 400   48, 40   40, 404     HEBBAIDOZEZ   52, 796   46, 044   90, 968   58, 504   47, 240   28, 411   37, 912   40, 048     HEBBAIDOZEZ   52, 796   46, 044   90, 968   58, 504   47, 240   28, 411   37, 912   40, 048     HEBBAIDOZEZ   53, 796   46, 044   90, 968   58, 504   47, 240   28, 241   37, 912   40, 048     HEBBAIDOZEZ   197, 859   27, 170   13, 313   16, 958   11, 1517   23, 904   29, 214   16, 599     HEBBAIDOZEZ   107, 859   24, 187   46, 646   22, 248   22, 590   37, 216   25, 827   21, 222     HEBBAIDOZEZ   107, 859   24, 187   46, 646   22, 248   22, 590   37, 216   25, 827   21, 222     HEBBAIDOZEZ   107, 859   24, 187   46, 646   22, 248   22, 590   37, 216   25, 827   21, 220     HEBBAIDOZEZ   107, 859   62, 187   586   57, 860   51, 172   48, 511   45, 193   47, 291     HEBBAIDOZEZ   107, 859   62, 187   59, 187   59, 187   59, 187   22, 492     HEBBAIDOZEZ   107, 859   62, 187   59, 187   59, 187   59, 187   22, 492     HEBBAIDOZEZ   13, 180   65, 241   15, 241   17, 241   23, 727     HEBBAIDOZEZ   17, 180   40, 718   73, 282   31, 403   30, 214   25, 214   23, 727     HEBBAIDOZEZ   17, 180   60, 718   73, 282   31, 403   30, 214   25, 214     HEBBAIDOZEZ   18, 181   89, 986   247, 534   84, 722   54, 176   46, 941   61, 944   77, 085     HEBBAIDOZEZ   18, 181   18, 183   65, 281   18, 182   19, 243   19, 244   19, 242     HEBBAIDOZEZ   18, 183   65, 18								8.869	5.823	12. 283
	5						71.673	113. 228	101.786	53. 514
							16.559		7.527	11.624
									42.964	19.680
Figure   1007-2515   195   59   197   207   104   68   171   48   643   58   647   41   135   59   659   107     Figure   107   68   77   78   78   78   78   78   7		,								40.048
										59, 667
TREEBATOWESS   107, 859   24, 187   45, 546   22, 248   22, 250   37, 216   25, 927   33, 282   TREEBATOWESS   33, 320   62, 916   259, 854   57, 860   53, 172   46, 511   45, 193   47, 291   TREEBATOWESS   33, 320   32, 914   154, 170   35, 196   22, 489   76, 280   22, 314   27, 272   14, 184, 184, 184, 184, 184, 184, 184,									29, 214	16.599
HEBBAI002785   333, 320   52, 916   259, 854   57, 860   51, 172   46, 511   45, 193   47, 291     HEBBAI002866   38, 358   58, 3014   45, 170   35, 158   22, 499   52, 290   22, 314   23, 727     HEBBAI002867   155, 384   165, 724   64, 658   79, 322   320, 013   30, 234   25, 512   83, 769     HEBBAI002873   71, 550   40, 718   73, 222   33, 403   39, 514   40, 129   38, 619   22, 532     HEBBAI002873   55, 486   61, 581   83, 386   247, 534   34, 722   54, 176   45, 941   51, 944   77, 085     HEBBAI002873   55, 486   61, 838   62, 47, 534   34, 722   54, 176   45, 941   51, 944   77, 085     HEBBAI002895   43, 813   17, 555   52, 241   14, 788   12, 463   31, 752   34, 100   18, 772     HEBBAI002895   43, 813   17, 555   22, 241   14, 788   12, 463   31, 752   34, 100   19, 772     HEBBAI002705   43, 533   30, 740   37, 573   54, 473   30, 588   113, 980   37, 727   59, 277     HEBBAI002705   43, 533   30, 740   37, 573   54, 473   30, 588   113, 980   38, 727   52, 277   22, 277   23, 224     HEBBAI002705   43, 533   30, 740   35, 579   18, 468   19, 118   26, 777   29, 277   27, 224     HEBBAI002705   43, 533   30, 740   35, 579   18, 468   19, 118   26, 777   29, 277   27, 224     HEBBAI002712   52, 678   59, 111   110, 506   41, 591   41, 592   43, 597   39, 504   30, 572   22, 54     HEBBAI002712   43, 404   51, 584   59, 685   41, 591   41, 41, 594   41, 597   41, 594     HEBBAI002712   43, 404   51, 585   77, 677   15, 783   23, 557   19, 684   27, 647   79, 70     HEBBAI002712   40, 404   51, 594   41, 404   51, 594   41, 404   51, 594     HEBBAI002713   77, 679   26, 461   34, 600   41, 594   41, 597   41, 41, 597   41, 41, 597     HEBBAI002734   77, 679   26, 461   34, 600   41, 598   41, 41, 597   51, 583   41, 400   41, 400     HEBBAI002730   31, 324   51, 585   51, 595   51, 500   51, 5	10									23, 282
HERBATODZEST   38, 495   88, 1014   154, 170   35, 195   22, 499   26, 290   27, 314   23, 727     HERBATODZEST   155, 384   165, 214   164, 656   29, 323   320, 013   30, 214   25, 612   38, 769     HERBATODZEST   155, 384   165, 214   164, 656   29, 323   320, 013   30, 214   25, 612   38, 769     HERBATODZEST   161, 681   89, 386   227, 314   44, 725   44, 61, 725   44, 61, 72   54, 768     HERBATODZEST   161, 681   89, 386   247, 534   44, 722   54, 176   45, 941   61, 944   77, 085     HERBATODZEST   161, 681   89, 386   247, 534   44, 722   54, 176   45, 941   61, 944   77, 085     HERBATODZEST   55, 416   61, 838   65, 537   37, 679   18, 172   29, 420   38, 238   44, 113     HERBATODZEST   36, 316   47, 735   54, 473   47, 473   47, 48, 48, 48, 48, 48, 48, 48, 48, 48, 48										
		***************************************							22.314	
HEBBAT002587   155.384   155.244   164.555   29.523   520.013   30.214   25.512   33.769     HEBBAT002578   161.681   89.986   247.534   84.725   54.176   45.941   61.944   77.065     HEBBAT002588   6.756   73.564   55.37   37.679   18.172   22.420   38.238   44.113     HEBBAT002588   6.756   73.564   55.37   37.679   18.172   22.420   38.238   44.113     HEBBAT002588   6.756   73.564   55.37   37.679   18.172   22.420   38.238   44.113     HEBBAT002589   49.633   17.555   29.241   14.788   17.93   4.605   3.500   14.777     HEBBAT002710   35.528   96.718   77.735   54.473   50.568   113.980   87.727   59.878     HEBBAT002712   52.578   59.111   101.506   41.591   41.597   35.579   35.643   30.722   28.457     HEBBAT002712   52.578   59.111   101.506   41.591   43.597   35.643   30.722   28.457     HEBBAT002712   52.578   59.111   101.506   41.591   43.597   35.643   30.722   28.457     HEBBAT002712   52.578   59.111   101.506   41.591   43.597   35.643   36.72   28.457     HEBBAT002712   52.578   59.111   101.506   41.578   43.597   35.643   36.72   28.457     HEBBAT002712   52.578   59.111   101.506   41.578   43.597   35.643   36.77   26.527     HEBBAT002712   52.578   59.111   101.506   41.578   43.597   35.643   36.77   26.527     HEBBAT002712   52.578   59.111   101.506   41.578   43.597   43.647   57.572     HEBBAT002712   71.798   88.550   29.109   81.299   43.679   55.807   56.822   37.603     HEBBAT002713   77.797   26.481   34.604   21.128   21.756   45.830   46.221   57.003     HEBBAT002730   70.678   70.848   83.500   29.309   81.598   43.679   55.802   8.2997   09.83   27.599     HEBBAT002730   70.678   72.803   35.400   15.803   15.500   15.803   43.499   31.400   49.691     HEBBAT002736   60.078   72.803   43.604   21.128   21.756   44.599   29.93   31.400   49.691     HEBBAT002736   70.756   72.505   72.072   41.798   31.291   54.649   39.349   39.737     HEBBAT002736   70.056   45.505   45.905   77.775   44.599   29.591   79.991   79.991   79.991   79.991   79.991   79.991   79.991   79.991										
HEBBAT002879										
HEBAT002579										22.532
### HEEBAT002785	15								61.944	
FIRESANDOZESS									38. 238	44.113
HEBBA10027703			1						3,600	2.944
HEEBA10027705								31.752	34, 100	14.772
FEEBA1002776									87.727	59.878
HEBRAT002715								26.777	29.277	29. 224
HEBBA10027718	20								30.872	26. 457
REPSA1002716					87.643		41.264	95. 279	127.808	65.580
#EFERA1002718							23.557			
				19.063	41.749	26.345		28. 367	26. 822	
### HEIBA1002773		HEMBA1002728	117. 984	88.950	293.019	81.290	43.679	65. 830		
HEBBA1002742 10.730 11.276 12.768 7.910 1.394 8.502 8.297 10.909 HEIBBA1002745 60.876 22.803 35.400 15.830 15.830 30.505 31.889 32.759 HEBBA1002745 76.748 26.130 38.669 17.760 32.813 43.493 51.440 49.651 HEBBA1002750 40.653 45.306 95.205 18.200 10.037 22.527 29.331 30.774 HEBBA1002755 94.758 62.505 220.964 63.414 37.572 44.593 28.497 39.737 HEBBA1002763 13.935 3117 8.450 3.792 2.291 8.714 10.261 5.285 HEBBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEBBA1002766 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEBBA1002766 103.210 30.236 54.998 17.099 19.753 35.636 41.922 26.940 HEBBA1002777 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEBBA1002777 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEBBA1002779 87.338 50.411 81.356 42.070 19.557 31.370 27.642 39.672 HEBBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002799 87.351 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.633 HEBBA1002805 5.941 88.407 13.385 50.2 25.574 33.389 50.974 49.045 HEBBA1002806 776 5.495 16.825 5.349 3.199 11.765 5.339 171.186 117.409 HEBBA1002806 77.5 5.945 84.076 13.829 84.939 37.555 59.859 54.302 55.958 HEBBA1002807 5.941 84.076 55.603 38.073 79.745 59.869 54.302 55.958 HEBBA1002806 5.941 84.076 55.603 38.073 39.502 25.574 33.389 50.974 49.045 HEBBA1002807 5.941 84.076 55.603 38.073 39.502 22.845 58.433 37.555 34.844 39.851 35.868 37.355 39.868 39.873 39.974 39.974 39.7755 39.974 39.974 39.7755 39.974 39.974 39.97		HEMBA1002730	131.726	26.862	67.877	28.628	36.686			
#EEBA10027745 60.876 22.803 35.400 15.830 15.630 30.605 31.889 32.759   #EEBA10027748 76.748 26.130 38.569 17.760 32.813 43.493 53.440 48.691   #EEBA1002755 40.563 45.306 95.205 18.200 10.037 22.527 29.331 30.774   #EEBA1002755 94.758 62.505 220.964 63.414 37.572 44.593 28.497 39.737   #EEBA1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285   #EEBA1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285   #EEBA1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347   #EEBA1002768 100.803 57.554 59.457 35.570 28.005 43.770 40.930 38.215   #EEBA1002770 103.210 30.236 54.098 17.099 19.753 35.566 41.922 26.940   #EEBA1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776   #EEBA1002770 79.7457 29.259 75.705 22.719 22.643 33.689 38.357 27.804   #EEBA1002770 79.7457 29.259 75.705 22.719 22.643 33.689 38.357 27.804   #EEBA1002780 77.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672   #EEBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672   #EEBA1002780 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761   #EEBA1002780 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761   #EEBA1002780 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761   #EEBA1002780 89.194 21.334 22.468 20.281 12.823 11.156 11.647 15.735   #EEBA1002801 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794   #EEBA1002801 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794   #EEBA1002801 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794   #EEBA1002806 55.084 37.823 55.994 35.902 25.574 33.389 50.974 49.045   #EEBA1002807 25.385 13.583 42.122 15.523 9.507 33.39 171.186 117.409   #EEBA1002805 5.941 8.407 13.251 6.179 2.332 4.352 4.364 3.363 57.345   #EEBA1002805 5.941 8.407 13.251 6.179 2.332 4.352 4.364 35.83	25	HEMBA 1002734	77.579	26.481	34.604	21.128				
HEEBA1002748 76.748 25.130 38.569 17.760 32.833 43.493 53.440 49.691 HEEBA1002750 40.663 45.306 95.205 18.200 10.037 22.527 29.331 30.774 HEEBA1002759 94.758 62.505 220.964 63.414 37.7572 44.593 28.497 39.737 HEEBA1002759 13.935 3.117 8.450 3.792 2.291 8.714 10.261 5.285 HEEBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEEBA1002766 100.803 97.554 59.457 35.570 28.006 43.770 40.930 38.215 HEEBA1002766 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEEBA1002777 20.350 16.288 28.054 21.736 10.754 12.030 14.991 11.776 HEEBA1002777 130.615 37.655 72.072 41.794 31.219 54.881 59.342 43.552 HEEBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEEBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.672 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEEBA1002790 87.371 61.291 152.514 38.033 1.755 41.334 100.167 80.301 50.035 HEEBA1002801 10.311 4.603 11.704 31.904 4.420 30.915 44.046 51.794 HEEBA1002801 10.311 4.603 11.704 31.904 4.420 30.915 44.046 51.794 HEEBA1002801 10.311 4.603 11.704 31.904 4.420 30.915 44.046 51.794 HEEBA1002806 76.504 77.515 97.755 97.		HEMBA1002742	10.730	11.276						
HEIBA1002750 40.663 45.306 95.205 18.200 10.037 22.527 29.331 30.774 HEIBA1002755 94.758 62.505 220.964 63.414 37.572 44.593 28.497 39.737 HEIBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEIBA1002767 65.682 25.272 35.782 14.035 19.183 31.497 33.393 18.347 HEIBA1002769 103.210 30.236 54.098 17.099 19.753 35.636 41.922 26.940 HEIBA1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEIBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 18.397 177.576 HEIBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 18.357 27.804 HEIBA1002780 72.338 50.411 181.356 42.070 19.957 31.370 27.642 39.6761 HEIBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEIBA1002799 97.457 29.259 75.705 22.719 22.643 33.689 18.357 27.804 HEIBA1002799 97.457 29.259 75.705 22.719 22.643 33.689 18.357 27.804 HEIBA1002799 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEIBA1002799 99.457 39.854 31.704 31.990 4.420 3.016 33.829 6.633 HEIBA1002899 9.194 21.334 22.468 20.251 12.823 11.156 11.647 15.735 HEIBA1002890 10.311 4.603 11.704 3.190 4.420 3.016 33.829 6.633 HEIBA1002810 42.533 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEIBA1002810 42.533 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEIBA1002810 42.533 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEIBA1002806 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEIBA1002807 139.924 107.278 533.137 90.533 79.745 99.869 54.302 52.988 HEIBA1002806 60.733 32.524 30.030 9.693 9.527 27.595 19.197 18.101 HEIBA1002806 47.76 6.495 16.825 5.349 3.119 11.765 7.385 8.35 117.409 HEIBA1002807 24.248 40.839 17.864 23.748 44.385 57.302 35.668 HEIBA1002806 75.834 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEIBA1002806 75.834 37.856 38.801 17.823 19.509 27.595 19.197 18.101 HEIBA1002806 75.834 37.856 38.801 17.803 19.500 24.356 55.374 45.005 HEIBA1002807 31.900 31.900 32.500 33.600 33.000 30.900 32.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.900 33.9			60.876	22.803						
HEBBA10027755				26.130						
HEBBA1002755		HEMBA 1002750	40.663							
HEBBA1002763 430.941 88.931 172.920 71.623 88.921 195.471 197.995 118.224 HEBBA1002766 100.803 57.554 59.457 35.570 28.006 43.770 40.930 38.215 HEBBA1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEBBA1002770 20.350 16.268 28.054 21.736 10.754 12.030 14.991 11.776 HEBBA1002770 30.351 57.655 72.072 41.794 31.219 54.881 59.342 43.552 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEBBA1002779 97.457 29.259 75.705 22.719 22.643 33.689 38.357 27.804 HEBBA1002790 87.371 61.291 152.514 38.033 29.616 28.032 20.352 34.761 HEBBA1002794 202.405 77.515 95.182 31.252 41.834 100.167 80.301 50.036 HEBBA1002798 9.194 21.334 22.468 20.281 12.823 11.155 11.647 15.735 HEBBA1002801 10.311 4.603 11.704 3.190 4.420 3.016 13.829 6.693 HEBBA1002816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEBBA1002816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEBBA1002816 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.404 HEBBA1002830 139.924 107.278 533.137 90.533 79.745 59.869 54.302 25.958 HEBBA1002826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEBBA1002836 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEBBA1002836 77.126 30.401 44.872 22.577 28.539 50.264 55.374 45.005 HEBBA1002836 77.126 30.401 44.872 22.577 28.539 50.264 55.374 45.005 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002836 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEBBA1002837 15.600 32.845 58.138 14.590 22.846 54.873 55.608 38.801 HEBBA1002931 65.007 59.109 104.125 15.411 19.920 31.099 23.998 19.182				62.505						
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HEMBAIO02810 42.583 45.313 55.088 35.416 29.480 60.935 44.046 51.794 HEMBAIO02816 52.084 37.823 56.994 35.902 25.574 33.389 50.974 49.045 HEMBAIO02818 321.516 100.826 187.799 84.893 81.695 152.339 171.186 117.409 HEMBAIO02826 40.776 6.495 16.825 5.349 3.319 11.765 7.355 8.363 HEMBAIO02833 119.102 44.248 40.839 17.864 23.748 44.398 57.302 36.668 HEMBAIO02850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02850 7.914 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBAIO02863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBAIO02867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBAIO02876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBAIO02886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBAIO02896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBAIO02896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBAIO02891 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBAIO02894 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBAIO02893 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889		HEMBA 10021 30								
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HEMBA1002850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735  HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101  HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005  HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196  HEMBA1002866 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689  HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574  HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662  HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801  HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182  HEMBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889								11.765		8. 363
HEMBA1002850 5.941 8.407 13.251 6.179 2.932 4.352 4.844 3.735 HEMBA1002862 60.735 32.524 30.030 9.693 9.527 27.595 19.397 18.101 HEMBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HEMBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEMBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEMBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEMBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HEMBA1002913 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HEMBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889										
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HETBA1002863 77.126 30.401 44.872 22.577 28.639 50.264 55.374 45.005 HETBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HETBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HETBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HETBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HETBA1002813 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HETBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HETBA1002934 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HETBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HETBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889										
HEEBA1002867 25.385 13.583 42.122 15.283 9.501 22.992 15.180 16.196 HEEBA1002876 101.249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HEEBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HEEBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.652 HEEBA1002813 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEEBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEEBA1002934 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEEBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HEEBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889							28. 539			
HETBA1002876 101,249 55.603 38.073 36.480 23.017 53.318 51.363 56.689 HETBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HETBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.652 HETBA1002891 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HETBA1002921 53.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HETBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HETBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HETBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889						15. 283				
FETBA1002886 9.474 14.188 23.688 7.657 11.980 14.640 6.432 18.574 HETBA1002896 78.580 27.420 49.774 16.754 20.366 36.684 35.283 42.662 HETBA1002891 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HETBA1002821 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HETBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HETBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HETBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889						36. 480				
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HEMBA1002813 126.001 32.845 58.138 14.590 22.846 54.873 56.608 38.801 HEMBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEMBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEMBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HEMBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889	50	HEMBA1002896								
HEPBA1002921 63.378 25.443 37.615 15.333 19.054 28.881 37.595 34.298 HEPBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEPBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HEPBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889				32.845	58.138					
HEEBA1002924 65.007 29.109 104.125 15.411 19.920 31.099 23.998 19.182 HEEBA1002934 432.841 308.291 644.522 180.470 145.293 273.733 166.153 242.809 HEEBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889				25. 443	37.615					
HEMBA1002935 92.005 52.184 221.722 49.477 41.867 34.331 29.646 38.889			65.007	29. 109						
(1000)100000   00.000   <u>-0.10.</u>   120.000		HELBA1002934	432.841							
55		HEMBA1002935	92.005	52. 184	221.722	49.477	41.867	34.331	29.646	38, 889
	55									

Table 24

	HEMBA1002937	38.698	30. 844	33.817	12.784	18. 251	14, 107	24. 131	18.662
					19.077	13.734	19. 266	17.364	17.750
	HEMBA1002939	39.755	22.867	33.838		18. 241		21.112	16. 286
5	HEMBA1002944	53. 762	33.349	51.861	21.860		23. 920		21, 177
	HEMBA1002951	38.716	29. 783	39. 196	19.808	29.614	19.702	28. 422	
	HEMBA1002954	24. 907	8. 542	20. 941	9. 265	13. 758	15.056	7.297	13. 424
	HEMBA1002962	86.680	62.578	220. 246	62.027	37. 753	44.037	31.812	41.725
	HEMBA1002968	105.871	78. 850	221.414	85.545	40. 380	43.093	38.816	50. 281
	HEMBA1002970	48. 034	34.741	30.834	18.482	6.639	17. 125	23.514	36. 180
40	HEMBA1002971	39. 492	44, 145	35.618	25.614	12. 932	25. 193	14.823	23. 202
10	HEMBA1002973	83.710	70.965	156, 167	43.307	28. 902	29.947	26. 101	34.769
	HEMBA1002978	35. 833	19. 362	27.056	13.075	20. 398	11.324	16.059	13.956
	HEMBA1002981	107. 112	35. 200	56.576	23.695	26. 105	33.054	37, 199	21. 249
	HEMBA1002985	79. 217	44. 154	116.532	27.950	26. 158	37.462	28. 927	20. 335
		61.056	78. 203	58.834	49.967	64. 529	38. 333	28.919	20. 529
	HEMBA1002986					8. 965	16.229	15.956	22, 796
15	HEMBA1002988	37.307	36.609	71.802	20.621	34. 289	57.004	61.291	91. 211
	HEMBA1002992	97.720	72.656	79.841	50.454			19.510	48. 529
	HEMBA1002995	51.473	63.779	55, 081	36.903	25. 007	38.630		25. 973
	HEMBA1002997	41.734	70.805	29. 264	27.019	33.664	24. 201	18.442	
	HEMBA1002999	35. 341	16.456	18. 357	11.146	7.034	12.086	13.966	9. 970
	HEMBA1003004	55. 654	33.689	35. 194	15.119	16, 204	20.866	27.891	20. 055
20	HEMBA1003006	40.682	24.886	20,750	20.903	26. 595	25. 445	20.310	20. 924
20	HEMBA1003008	29.269	20. 922	74.697	25.061	17. 787	10. 271	5. 688	12.638
	HEMBA1003021	130.889	123.646	311.225	101.957	95. 443	64.844	60.969	90. 296 -
	HEMBA1003027	54. 935	32.610	44.710	18.890	52. 131	26.286	28.112	31.561
	HEMBA1003029	33. 333	42.436	60.787	20.829	34. 111	29.704	49. 230	45. 833
	HEMBA1003031	34.000	25. 311	18.494	14.998	13. 316	13.955	15.773	27.136
	HEMBA1003032	171, 114	46.990	71.365	23.640	50. 526	81.278	84.036	46. 352
25	HEMBA1003033	168.563	118.674	378.771	109. 222	90.670	70.150	55. 336	77.819
	HEMBA1003034	173, 162	127.221	484, 135	108.238	85. 630	61.733	36.799	63. 312
	HEMBA1003035	11.693	5. 195	9.305	4.478	5. 058	11.024	2. 553	4. 409
	HEMBA1003037	261.159	89. 481	145, 321	58. 521	65.732	104.677	89.571	71. 674
	HEMBA1003041	103.945	105.085	291.931	93. 188	75. 193	53.097	39.564	58. 217
	HEMBA1003046	40. 254	39.965	46.856	26.192	11.615	35.659	25. 378	32.416
30	HEMBA1003047	127.888	49. 341	139.750	32.219	32. 320	57.450	33.390	28.702
	HEMBA1003048	87.433	35. 962	42.305	12.040	20.442	39. 108	29.597	21.461
		6.366	8. 535	6.201	8.809	4. 415	7. 239	3.330	7.829
	HEMBA1003064		34.508	77.097	26. 154	20. 523	28.755	24.783	17. 488
	HEMBA1003067	55.833		28.869	17.461	19.647	20.624	22.285	19. 438
	HEMBA1003071	54.728	22.509	31.225	26. 146	22.906	21.483	17.616	19. 134
	HEMBA1003072	62.421	30.769			40.422	49.911	47.023	64.737
35	HEMBA1003076	111.254	51.085	78. 972	37.151	8. 453	18.661	13. 797	5.837
	HEMBA1003077	36.471	15. 407	24. 522	8.009	37.169	17. 933	17. 439	18. 923
	HEMBA1003078	34. 143	38.741	77. 906	31.907		25. 576	18.026	24. 526
	HEMBA1003079	28. 559	39.563	41.646	26.110	25. 889	41.611	29.619	67.469
	HEMBA1003083	61.036	48. 635	169.439	52.788	60.016		16.039	19. 219
	HEMBA1003086	49.032	40.488	154. 409	29.869	12.063	16.544	25. 027	14. 265
40	HEMBA1003090	34.778	14.860	23.758	12.710	24. 132	15.848	84. 551	34.775
	HEMBA1003094	184. 999	43. 363	72.116	30.096	53.636	78. 251		11.348
	HEMBA1003096	31.440	18.030	25.774	10.290	11.781	14.033	27.791 18.003	31.059
	HEMBA1003098	36.774	64. 970	88. 562	34.074	24. 271	25.656		
	HEMBA1003101	55.716	24. 121	22.316	11.682	13. 163	21.315	25. 117	15. 689
	HEMBA1003109	48.411	21.093	39.285	21.315	21.724	27.826	31.034	21.809
45	HEMBA1003114	41.101	24.786	22.792	14. 164	14.657	18.320	15. 152	16.038
45	HEMBA1003117	22.939	13.535	20, 191	6.812	10. 538	14.917	18.015	12.566
	RELIBA1003120	24.531	24. 408	55. 805	26. 574	13.838	15.423	15.080	21.728
	HEMBA1003129	40.276	46.792	104.463	37.995	37. 989	21.990	26. 267	38. 207
	HEMBA1003133	50.080	22.873	35.022	15.164	20.000	21.592	25.551	27.656
	HEMBA1003136	146.630	23.706	65.990	18.301	31.049	69.754	51.669	25. 346
	HEMBA1003142	69.008	47.867	130.557	32.955	30.384	25.274	27.118	29. 493
50	HEMBA 1003 148	59.282	20.084	32.740	18. 292	18. 973	32.206	22.003	24.674
	HEPBA1003151	53.856	20.003	51.824	13. 233	9.854	27.114		13.546
	HEMBA1003152	20.577	9. 803	19.388	10.017	5.761	31.586	23.227	6.853
	HEMBA1003157	16.477	9. 272	16. 246	9, 919	17.605	7.547		10. 181
	HEMBA1003166		257. 380	671.361	260, 521	221. 325			
		293.814				6.963	7.361	5.733	
55	HEMBA1003171	17.730	8. 702	16. 527	6.499	1 0.303	1.301	3.700	
-									

Table 25

					-00 FO. T	17 634 1	A1 150 T	13 300	15 025
	HEMBA1003175	38.620	40.445	100. 302	29. 594	17.624	21.152	13. 386	15. 936
	HEMBA1003179	63.835	33.869	50.631	27. 163	25. 502	35, 500	39.052	37.713
	HEMBA1003186	100.461	75.611	231.787	75. 781	58.278	54. 222	55.862	61.615
5									
	HEMBA1003196	36.422	27.557	45. 633	20.623	18.740	21.756	30. 501	35.864
	HEMBA1003197	8.462	9. 564	5. 534	5. 965	4. 051	3.138	7.054	7.066
	HEMBA1003199	34.650	18. 409	81. 183	15.696	16.799	9. 492	17.381	15. 917
	HEMBA1003202	79.337	59.764	236.822	43.286	41.820	31.106	32, 936	45. 183
	HEMBA1003204	66.523	56. 272	172.818	48.560	31.451	33.193	27.421	28.849
	HEMBA1003210	23.713	52.768	35. 498	5. 529	38.451	16.353	59.417	17.563
10							59.663		54. 363
	HEMBA1003212	126.394	90.709	372.474	74. 164	62.392		45.714	
	HEMBA1003218	19, 415	13, 105	13.670	6.371	4,792	13.681	10.789	6. 536
	HEMBA1003220	81, 171	86.642	147.453	89, 495	42.391	47.586	54.647	123.019
							10.413	9.856	14. 985
	HEMBA1003222	25.803	22. 891	28. 577	7.994	11.404			
	HEMBA1003225	105.735	21. 238	40.848	11.586	20.280	48. 243	44.574	19. 547
	HEMBA1003229	30.394	26, 363	41, 333	22.998	17.475	14.707	20. 154	19.749
15					31.176	20,775	56.815	40, 191	75, 238
	HEMBA1003230	69.643	70.015	42.439					
	HEMBA1003235	44. 989	43.337	105. 267	33.038	19. 405	20.834	22.018	29.856
	HEMBA1003236	8.677	17.896	8.735	7.270	7. 328	17. 286	5. 295	18.441
	HEMBA1003250	7. 260	12. 598;	12.993	4.750	4.815	7.242	5. 982	4. 378
									60.076
	HEMBA1003252	56.274	51, 495	65. 197	28. 241	33.512	44.917	62.506	
	HEMBA1003257	71.751	16.083	40.414	13.391	19.441	38. 988	28.614	19.028
20	HEMBA1003268	19.492	18, 996	46. 948	14, 167	12.769	11.524	8. 522	17.414
			38. 933	125. 242	29.404	21, 135	22. 989	17. 240	24, 704
	HEMBA1003273	48. 113							26, 145
	HEMBA1003276	36.279	34, 802	113. 584	23.812	17.208	20. 437	14.685	
	HEL'8A1003277	31.363	12.827	21.514	10.462	11. 287	13. 206	16. 182	14. 465
	HEI'BA1003278	36.998	24.906	71.222	17.479	15.791	16.787	10.948	17. 841
				38.057	16.933	20, 792	37. 901	30.931	31. 493
25	HELBA1003280	50.716	16.000						
25	HEYBA1003281	66.732	21. 393	32.728	15.032	18.415	26.844	28. 577	24. 898
•	HEMBA1003284	9.746	8. 482	12.941	5.779	5.747	5.813	3. 545	3.499
	HEMBA1003286	69.502	35. 947	60.729	21.827	29.473	52.233	50. 283	47.695
						18, 100	6. 561	6.341	7,647
	HEMBA1003291	13. 248	9, 951	10.909	3.504				
	HEMBA1003294	69. 599	52.239	168.555	39. 127	38.460	40.377	24.057	27. 486
	HEMBA1003296	61.933	31.456	37.947	21.206	23, 199	23.249	34. 580	37.768
30	HEMBA1003304	7, 117	5. 972	8. 976	6.154	8.839	4. 199	3, 461	3. 227
00								6. 333	9. 387
	HEMBA1003306	17. 590	15. 590	22.443	8.410	11. 282	8. 448		
	HEMBA1003309	6. 845	10.103	12.198	14.015	7,776	8.709	3. 955	18. 326
	HEMBA1003314	637.052	210, 608	238.618	105.098	198, 106	299.884	273.738	171.516
				84.690	32.381	29.482	56.694	53. 105	54. 024
	HEMBA1003315	83.736	51.612						
	HEMBA1003322	108.401	88.539	256.570	51.502	51.083	44. 130	42.804	45. 519
35	HEMBA1003326	42.723	20, 581	14.759	11.799	7.780	18. 087	12.420	9. 516
	HELBA1003327	61.811	36.702	87.698	28. 181	19.784	18. 596	17. 453	18. 377
						25.000	18.669	22.079	32.865
	HEMBA1003328	53. 406	51.712	114.941	36.926				
	HEMBA1003330	108. 955	82.099	207.708	73.413	52.244	50.838	55. 920	55. 390
	HEMBA1003348	121.625	110.275	337.182	94, 209	99.717	67.000	43.513	80.023
	HEMBA1003369	5. 861	23.644	14.930	4.979	1.726	9.064	3,020	5. 373
							140.758	150.458	124, 948
40	HEMBA1003370	315.016	197.956	369.117	140.044	139.216			
	HEMBA1003373	50. 135	31.291	53.330	17.430	5.513	19.164	8.117	19, 638
	HEMBA1003376	174. 269	170.290	519.668	126.099	89.798	108. 226	81.818	107.084
	HEMBA1003380	43.015	24.657	74.071	29.281	24, 407	19,711	13.485	20,047
				68.079	15. 389	9. 455	11.810	8.800	14. 281
•	HEMBA1003384	25, 555	30.071					<del></del>	
	HE38A1003387	6.515	2.588	2.697	1.577	1.109	1.803	1.986	3.464
	HEMBA1003392	111. 457	25.882	42.253	17. 323	29.007	50.086	29.337	23.550
45		16.068	18,666	35.483	15.254	9.873	10. 355	6, 207	12.514
	HEMBA 1003395							27.860	34.116
	HEMBA1003399	45. 227	21.480	37.035	19.231	15. 354	19.471		
	HEMBA1003400	116, 210	36.907	58.706	24.811	49. 133	53.819	60.041	53. 109
	HEMBA1003402	32.500	16.239	27.854	8.795	12.867	17.141	11.617	14, 596
								44.891	45. 870
	HEMBA1003403	60.260	43.377	46.720	20, 221	26. 579	36.738		
	HEMBA1003408	196.676	49.687	70.460	29.354	50.910	84. 358	77.062	45. 433
50	HEMBA1003412	104. 813	43.934	55.699		43.763	61.953	59.463	47.139
				25.036		7. 282	10.593	5.696	11.032
	HEYBA1003417	22. 445	13.970						
	HEMBA1003418	57. 411	57.397	76.232		45. 336	43.450	22.206	90, 604
	HEMBA1003420	29, 838	15, 856	201.831	11.319	8.067	11.379	12.938	14, 721
	HEMBA1003425	17, 466	15.895	21.662		6.723	8.483	10, 838	9, 083
									11. 581
55	HEMBA1003433	23, 931	18. 435	24. 576	12.136	10. 421	10.074	11.092	11. 361
55							_		

Table 26

									36 646
	HEMBA1003440	91.727	41.727	39. 257	19.755	26.941	45.998	31.620	35.845
	HEMBA1003442	7.090	22. 535	10, 452	33.897	10. 259	15.118	7.093	14.790
						18. 587		42.314	35.065
5	HEMBA 1003447	82. 161	36.670	48. 248	26.789		41.591		
3	HEMBA1003453	50, 472	26.692	25. 954	16. 130	11. 252	16, 584	28. 534	21.256
		55. 687	25. 328	42.686	17. 261	18.856	27.281	22.795	17.854
	HEMBA 1003461								
	HEMBA 1003463	40. 102	23.311	34. 469	13.456	19.704	20.277	16.984	18.124
	HEMBA 1003465	92, 245	40.963	61.816	28.410	36.051	39, 389	40.220	36. 851
								51.589	62.191
	HEMBA 1003480	114.075	114.841	266.076	76. 366	67.942	56.459		
	HEMBA 1003485	44. 403	28. 836	33.659	14. 371	8.636	26.284	16.036	14. 582
10	HEMBA1003487	42.939	15. 463 .	23.730	9,752	15.729	24.902	21.136	16. 494
	HEMBA 1003492	31.026	21. 538	56.674	14. 934	12.014	12.082	9. 567	14.655
	HEMBA 1003494	97. 366	260. 496	50.174	48. 821	12, 504	74.554	20.623	180.841
						13.881	21.376	18.562	6.072
	HEMBA1003497	39.000	17. 943	24. 659	11.432				
	HEYBA1003503	54.774	21.486	28. 175	12.948	17, 154	30.911	36.453	16. 806
	HEMBA1003511	18.672	14,740	43.023	11.794	13.330	8.925	16.405	11.615
15								179, 430	96.672
75	HEMBA1003528	385. 123	191.234	239.319	81.329	123.915	213.945		
	HEMBA1003530	43.820	12.384	23.693	10, 695	21.216	20.067	28.030	16.204
	HEMBA 1003531	111, 104	73. 542	215.578	67, 833	214.022	56. 139	50.217	66. 992
	HEMBA1003532	145. 137	62.379,	83.827	37. 506	53. 388	90.314	77.728	60.515
	HEMBA1003538	61.123	20.746	32.949	11.160	19. 286	34.305	28. 231	13.837
					5. 904	7, 197	10.239	6.617	8. 168
	HEMBA 1003545	21.489	10.501	20.608					
20	HEMBA 1 003546	31.371	32. 365	28.613	13.365	226.243	16.427	16.554	24.821
	HEMBA 1003548	4. 456	8. 124	9.845	4.563	7.542	6.155	5.647	8387
							49. 154	63. 525	48. 955
	HEMBA1003553	79.837	51.515	50.379	23. 327	28. 564			
	HEMBA 1003555	20.066	8.873	13.692	4.762	3. 584	10.112	10.962	6. 521
	HEMBA1003556	57. 280	36.399	128.391	29. 283	16, 426	19.257	18. 121	24.622
							2.983	6.207	6, 539
	HEMBA1003560	9. 290	4.426	2.529	2.848	1.767			
25	HEMBA 1003565	42.648	29. 588	20.996	8.344	13, 984	21.927	21.847	22.043
	HEMBA1003568	7.244	1.649	7.712	2, 430	3.763	3. 172	2.836	2, 592
							16.235	19.512	16, 518
	HEMBA 1003569	25. 048	20. 536	23.764	33. 957	13.740			
	HETALA 1003571	111.721	94.378	326.335	84. 368	71.788	50.029	48.011	59, 960
	HEMBA 1003579	3. 335	7. 399	15.353	6.553	8. 948	2.872	9.198	5. 421
					46.606		100 042		
	HEMBA1003580	274. 105	50. 292	102. 103	26.686	59.875	128.943	110.375	35. 695
30	HEMBA1003581	112.013	31.295	94.083	21.641	36.215	54. 336	50.711	21. 238
30			64.326	77.160	89. 876	47.882	53.615	40.656	45. 172
	HEMBA 1003591	97.076							
	HEMBA1003595	32.697	22.842	84.629	19.075	11.339	6.305	5. 581	18.085
	HEMBA1003597	48, 561	25.846	108.491	20.931	15.952	19.375	17. 580	20. 153
				22.468	12.142	11.688	18.934	21.743	15.025
	HEMBA1003598	49.728	20.134						
	HEMBA 1003600	32.772	35.099	56.905	26.268	29. 290	38.873	53, 305	56.783
	HEPBA1003602	18. 248	10.116	16.162	6.182	10.970	8.064	14.736	17.188
35			53. 579	69.723	24. 549	49. 902	105. 181	98, 166	47. 144
	HEMBA1003604	205. 949							
	HEMBA 1003610	140.996	29. 255	95.048	15. 492	103.150	72. 233	54.670	30,688
	HEMBA1003615	57.258	20.035	34. 102	12.808	16.022	24. 378	18.759	20.876
	HEMBA1003617		20.375	29, 789	12.148	22. 291	18. 199	18.770	18, 242
		48.414							
	HEMBA1003620	52.899	22.318	45.502	19.575	19.962	25. 239	39.072	29, 451
	HEMBA1003621	102.827	102.094	226.373	80.194	64.742	58.874	67.142	60.680
40	HEMBA1003622	19.815	13.838	25.009	16.055	8. 339	12.261	15.369	13.833
								10.303	11.422
	HEMBA1003630	20.008	16.381	30.244	13.871	5. 573	9. 992		
	HELBA 1003637	37.880	29.848	106.379	23. 251	18.468	17. 181	12.409	18. 500
	HEMBA 1003640	39.068	31.672	100, 901	22.572	22. 223	21.513	17, 417	20. 420
									5,711
	HEMBA1003645	25. 820	19.380	48. 445	13. 481	9. 247	12.142	54. 230	
	HEMBA1003646	38. 243	16.329	22.003	9.624	13.311	24.606	19.177	19.938
45			10.718	12.323	7.860	7.892	7.607	7.882	10,058
,,,	HEMBA 1003647	10.261							
	HEMBA 1 003656	40.171	31.269	66, 874	28. 981	19.429	18.898	23. 172	30. 178
	HEMBA1003662	25. 325	17.011	19.352	6. 387	10.041	10.909	14.055	18. 544
							9.774	15. 332	13.851
	HEMBA 1003666	23.086	11.187	17.407	5.803	8. 262			
	HEMBA 1003667	304.975	209. 929	337. 134	96.636	131.792	179.317	140.769	174. 256
	HEMBA 1003670	12.944	8.894	15. 235	3.344	2.565	7.057	6. 425	7.073
50							66.675	65. 424	47.173
50	HEM3A1003674	143.262	32. 196	51.919	33.863	62.734			
	HEYBA1003677	80.516	45. 946	220.695	45. 985	43.474	38.916	30. 594	46.808
	HEMBA 1003679	25. 325	7.795	16.167	6.727	5, 941	12.433	12.034	11.720
								38.990	39.343
	HEMBA1003680	42.317	25.723	33.794	24.664	23. 985	25, 419		
	HEMBA1003684	18.273	10.175	17.733	13. 315	4. 937	9.099	10. 182	10.574
	HEMBA1003690		65.531	75.876	46. 324	43.039	71,797	85.431	56.592
	UEYDA I NO 2020	115.021	1 03.331	13.010	1 40. 324	73.033	1 11.131	1 00. 701	1
55									

Table 27

HEBBA1003714   53, 732   57, 803   78, 875   73, 703   73, 703   74, 875   75, 875									
HEIBAN 1003702	HEMBA1003692	83, 253	96, 347	194, 372	63.188	40.872	47.354	39. 288	42.644
	HEMBA 1003702				23, 719	29,023	42.879	46,956	36.550
HEBBA 1003714									36.807
HEBBA 10037173   70   551   38   747   170   572   38   31   122   28   832   20   540   26   78   78   78   78   78   78   78   7									
HEBBAI003772	<u> </u>								
HEBBA10037729   83, 187   94, 829   133, 285   53, 886   49, 519   43, 330   22, 099   41, 184   184									
HEEBAI003778									
HEEBAI003778	HEMBA1003720	83. 587	94.829	133. 285					
HEIBAID03773	HEMBA 1003725	46. 157	55. 932	71.704	30.085	21. 305			31.573
HEEBAI003743	HEMBA1003728	103, 795	35, 668	58. 184	16. 485	21.818	42. 286	37.790	34. 280
HEBBA1003742				47,663	20, 231	15.376	18. 567	21.294	17. 427
HEEBA1003743   52. 409   32. 781   76. 684   22. 919   83. 476   18. 921   13. 867   14. 22   HEEBA1003743   75. 518   22. 518   23. 527   72. 589   25. 589   25. 581   42. 527   44. 100   24. 86   44.							3.652	3,024	4. 336
HEEBA1003742   40. 426   20. 265   50. 667   25. 589   21. 518   42. 057   44. 130   24. 86   HEEBA1003743   25. 518   27. 18   23. 392   18. 885   18. 530   12. 536   34. 868   73. 42   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 43   46. 2017   73. 44   46. 2017   73. 45									14. 220
HEEBA1003743	NEDBA1003733								
HEBBA1003760									
HEIBAIO03764 45.855 30.330 82.720 23.891 19.630 164.051 37.797 57.86 HEIBAIO03773 67.58 47.27 62.42 27.134 32.047 46.499 39.296 38.9 HEIBAIO03773 67.58 47.27 62.42 27.132 12.002 9.850 33.994 29.817 13.11 HEIBAIO03783 17.751 16.975 23.942 16.465 13.884 6.842 9.757 20.65 HEIBAIO03784 13.00 17.233 21.849 13.856 12.436 17.394 11.099 13.1 HEIBAIO03784 36.642 303.008 322.299 109.371 145.316 285.778 287.77 20.65 17.20 17.23 17.20 17.23 17.20 17.23 17.20 17.23 17.20 17.23 17.20 17.23 17.20 17.23 17.24									
HEIBAI003769   87.589   47.227   62.942   27.144   32.047   46.499   39.295   38.99   HEIBAI003773   37.575   16.975   23.942   17.002   9.850   33.904   29.817   13.11   14.1027   13.11   14.1027   13.11   14.1027   13.103734   13.604   13.500   17.233   21.849   13.856   12.436   17.334   11.095   13.11   14.1027   13.10374   38.64   42.303   22.299   10.9371   145.316   286.778   287.377   239.94   14.103794   38.64   33.003   22.299   10.9371   145.316   286.778   287.377   239.94   14.1030303   33.94   23.099   29.603   15.022   13.775   16.550   24.428   19.44   14.1030303   23.548   21.899   44.323   20.132   18.580   28.795   24.744   35.94   14.1030303   23.548   21.899   44.323   20.132   18.580   28.795   24.744   35.94   14.1030303   23.648   23.899   44.323   20.132   18.580   28.795   24.744   35.94   14.1030303   23.649   23.103	HEMBA 1003760			26.318					
HEEBA1003783	HEMBA1003764		30. 390						
HEIBRI 1003783 17.751 16.975 23.942 16.465 13.884 6.842 9.757 20.65 HEIBRI 1003784 13.500 17.233 21.849 13.556 12.436 17.394 11.099 13.1 HEIBRI 1003794 38.642 303.008 322.299 109.371 145.316 285.778 287.377 239.9 HEIBRI 1003793 39.392 23.099 29.603 15.022 13.775 16.550 24.428 19.4 HEIBRI 1003803 33.548 21.899 44.323 20.132 18.580 28.795 24.428 19.4 HEIBRI 1003804 80.382 26.816 48.558 16.154 27.867 31.087 37.611 22.5 HEIBRI 1003805 103.669 42.485 42.930 19.9371 145.516 288.795 24.744 35.9 HEIBRI 1003807 17.77 13.940 25.512 9.492 6.870 37.7 43.797 32.147 28.3 HEIBRI 1003807 21.777 13.940 25.512 9.492 6.870 37.7 43.797 32.147 28.5 HEIBRI 1003807 21.777 13.940 25.512 9.492 6.870 39.649 8.812 7.6 HEIBRI 1003807 21.777 13.940 25.512 9.492 16.870 37.611 22.5 HEIBRI 1003807 27.177 13.940 25.512 9.492 15.476 20.338 17.855 7.640 4.451 6.5 HEIBRI 1003807 27.777 29.7 42.7 42.5 42.9 42.9 11.5 47.6 HEIBRI 1003807 27.7 13.9 40 25.512 9.492 15.5416 219.584 26.07 27.83204 241.1 HEIBRI 1003808 27.3 42.9 42.4 21.1 HEIBRI 1003808 27.3 42.9 42.4 21.1 HEIBRI 1003808 27.3 42.9 42.4 42.1 11.4 HEIBRI 1003808 27.3 42.9 42.4 42.3 42.3 42.3 42.3 42.3 42.3 42.3	HEMBA1003769	87.589	47.227	62. 942	27.144	32.047			38. 944
HEIBA1003784	HEMBA1003773	63.842	14.722	21.132	12.002	9.850	33. 904	29.817	13.165
HEIBA1003794						13.884			20.650
HEEBA1003798   386.642   303.008   322.299   109.371   145.316   286.778   287.377   239.99   HEEBA1003799   39.395   23.099   29.603   15.022   13.775   16.550   24.428   19.4   19.							17.394	11.099	13.140
HEPBA1003893   39.392   23.099   29.603   15.022   13.775   16.550   24.428   19.44									239. 938
HEBA1003803   63.548   21.899   44.323   20.132   18.580   28.795   24.744   35.9     HEBA1003804   80.332   26.816   48.558   16.154   27.867   31.087   37.611   22.6     HEBA1003807   21.717   13.940   25.512   9.492   6.870   9.649   8.812   7.6     HEBA1003810   20.102   11.572   7.558   20.338   17.855   7.640   4.451   6.5     HEBA1003827   432.954   219.520   240.291   155.416   219.584   266.037   283.204   241.1     HEBA1003338   177.311   35.831   482.334   146.466   136.063   93.790   92.728   122.2     HEBA1003338   223.674   185.295   641.368   134.002   79.993   115.711   87.137   118.9     HEBA1003343   13.867   10.178   27.409   17.850   21.104   13.382   11.701   13.6     HEBA1003364   133.994   57.556   58.738   34.962   50.550   55.395   40.861     HEBA1003365   70.198   95.527   253.525   75.110   52.628   51.958   45.837   48.8     HEBA1003366   95.701   54.991   58.921   15.44   18.103   99.421   48.8     HEBA1003366   95.701   54.991   58.923   31.090   16.810   92.271   16.7     HEBA1003366   95.701   54.991   58.923   31.090   14.733   69.451   48.174   43.4     HEBA1003386   95.701   54.991   58.923   31.090   41.733   69.451   48.174   43.4     HEBA1003386   77.675   69.096   172.686   55.895   47.956   67.668   44.488   30.5     HEBA1003386   77.675   69.096   172.686   55.189   57.291   31.275   22.8     HEBA1003386   77.675   69.096   172.686   55.189   57.291   31.275   22.8     HEBA1003386   77.675   69.096   172.686   55.189   57.295   67.356   41.499   57.256   67.568   44.488   51.8     HEBA1003389   12.753   8.056   15.506   7.762   16.057   31.271   387.408   51.5     HEBA1003389   12.753   8.056   15.506   7.762   16.057   31.271   367.408   51.5     HEBA1003389   23.775   21.109   26.629   10.006   23.309   14.294   24.44   31.9     HEBA1003389   37.82   39.491   11.984   20.97   23.509   16.793   14.124   20.4     HEBA1003393   28.656   8.73   15.689   7.298   15.499   57.256   63.551   68.8     HEBA1003395   31.636   253.614   316.882   31.879   31.5429   63.070   22.3	the state of the s								19.403
HETBA1003804   80.382   26.816   48.558   16.154   27.687   31.087   37.611   22.6     HETBA1003805   103.669   42.485   42.930   19.984   36.377   43.797   32.147   28.3     HETBA1003807   21.717   13.940   25.512   9.482   6.370   9.649   8.812   7.6     HETBA1003827   432.964   219.520   240.291   155.415   219.584   266.037   283.204   241.1     HETBA1003836   177.311   135.831   482.334   146.465   136.063   93.790   92.728   122.2     HETBA1003837   177.311   135.831   482.334   146.465   136.063   93.790   92.728   122.2     HETBA1003838   223.674   185.295   641.368   134.002   79.993   115.711   87.137   118.9     HETBA1003843   13.867   10.178   27.409   17.850   21.104   13.382   11.701   13.6     HETBA1003846   133.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2     HETBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.5     HETBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.5     HETBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.5     HETBA1003864   52.130   18.071   24.567   9.568   13.009   16.810   29.271   16.7     HETBA1003864   52.730   18.071   24.567   9.568   13.009   16.810   29.271   16.7     HETBA1003864   52.730   18.071   24.567   9.568   13.009   16.810   29.271   16.7     HETBA1003867   62.950   44.572   159.217   48.098   42.466   37.097   36.010   45.8     HETBA1003867   60.003   60.0									35.938
HEUBA1003805									22.634
HEPBA1003807   21.717   13.940   25.512   9.492   6.870   9.649   8.812   7.6     HEPBA1003807   22.102   11.572   7.558   20.338   17.855   7.640   4.851   6.5     HEPBA1003827   42.964   21.950   24.0291   155.416   21.9584   266.037   28.3.204   241.1     HEPBA1003836   177.311   135.831   482.334   146.466   136.063   93.790   92.728   122.2     HEPBA1003838   223.674   185.295   641.368   134.002   79.993   115.711   87.711   18.73   118.9     HEPBA1003843   33.867   10.178   27.409   77.850   21.104   33.382   11.701   13.6     HEPBA1003846   133.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2     HEPBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   39.9     HEPBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   39.9     HEPBA1003856   27.277   12.805   22.440   12.068   13.009   16.810   29.271   16.7     HEPBA1003864   52.130   18.071   24.567   9.568   13.009   16.810   29.271   16.7     HEPBA1003869   59.701   54.991   58.923   31.090   41.733   69.461   48.174   43.4     HEPBA1003880   134.462   70.074   103.271   50.699   47.956   57.668   44.498   30.5     HEPBA1003881   134.462   70.074   103.271   50.699   47.956   57.668   44.498   30.5     HEPBA1003885   77.675   69.096   172.968   55.129   49.424   41.309   24.247   31.5     HEPBA1003893   386.525   281.955   515.307   187.300   180.355   21.81   29.281   31.275   22.8     HEPBA1003893   386.525   281.955   515.307   187.300   180.355   21.964   317.277   22.3     HEPBA1003893   386.525   281.955   515.307   187.300   180.355   21.964   317.277   22.3     HEPBA1003894   37.755   21.109   46.629   17.368   55.500   67.765   69.096   172.968   55.181   29.281   31.275   22.8     HEPBA1003895   77.675   69.096   172.968   55.500   55.000   50.385   21.81   29.281   31.275   22.8     HEPBA1003896   411.418   232.899   382.182   144.104   165.806   233.857   186.700   143.5     HEPBA1003895   77.675   69.096   172.968   55.500   50.378   19.070   14.294   24.342   17.68   14.294   24.34									
HEPBA1003810   20.102   11.572   7.558   20.338   17.855   7.640   4.451   6.55     HEPBA1003827   432.984   219.520   240.291   155.416   219.584   266.037   283.204   241.1     HEPBA1003838   27.674   185.295   641.368   34.002   79.993   115.711   87.137   118.9     HEPBA1003838   23.674   185.295   641.368   34.002   79.993   115.711   87.137   118.9     HEPBA1003843   13.867   10.178   27.409   17.850   21.104   13.82   11.701   13.6     HEPBA1003864   13.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2     HEPBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.9     HEPBA1003856   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.9     HEPBA1003865   27.378   13.868   16.982   14.248   8.662   11.259   9.145   9.9     HEPBA1003866   27.257   12.805   22.440   12.068   15.414   19.103   9.2271   16.7     HEPBA1003866   27.257   12.805   22.440   12.068   15.414   19.103   9.229   7.5     HEPBA1003866   27.257   12.805   22.440   12.068   15.414   19.103   9.229   7.5     HEPBA1003867   62.950   44.572   159.217   48.098   42.446   37.097   36.010   45.8     HEPBA1003868   95.701   54.991   58.923   31.090   41.733   69.461   48.174   43.4     HEPBA1003867   60.203   22.185   33.585   15.896   24.346   37.097   36.010   45.8     HEPBA1003867   76.767   69.096   17.968   55.129   49.424   41.309   24.247   31.5     HEPBA1003897   60.203   22.185   33.582   16.896   21.181   29.281   31.275   22.8     HEPBA1003896   411.418   232.899   38.185   16.896   21.181   29.281   31.275   22.8     HEPBA1003890   38.6525   281.955   515.307   187.300   180.355   212.964   37.297   122.3     HEPBA1003890   37.323   39.491   11.994   20.297   23.509   16.793   34.124   20.4     HEPBA1003890   37.323   39.491   11.994   20.297   23.509   16.793   31.1775   22.8     HEPBA1003890   37.323   39.491   11.984   20.297   23.509   16.793   31.1775   22.8     HEPBA1003890   37.605   33.857   38.656   37.755   20.006   33.1879   34.124   20.4     HEPBA1003891   37.905   63.862									
Heighton									
HEEBA1003836			11,572						
HEEBA1003838   223.674   185.295   641.368   134.002   79.993   115.711   87.137   118.9   HEEBA1003843   13.867   10.178   27.409   17.850   21.104   33.82   11.701   13.6   HEEBA1003846   133.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2	HEMBA1003827	432. 964							
HEEBA1003843	HEPBA1003836		135.831						
RESBA   D03846   133.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2	HEMBA1003838	223.674	185. 295	641.368	134.002	79. 993			118.957
HETBA1003846   133.994   57.556   58.738   34.962   50.550   56.395   40.861   60.2	HE™8A1003843	13, 867	10.178	27. 409	17.850	21.104	13. 382		13.634
HEMBA1003856   27, 378   13, 868   16, 982   14, 248   8, 662   11, 259   9, 145   9, 9		133, 994		58, 738	34, 962	50.550	56. 395	40.861	60. 253
HETBA1003857 101.908 95.527 253.525 75.110 52.628 51.958 45.837 48.8 HETBA1003864 52.130 18.071 24.567 9.568 13.009 16.810 29.271 16.7 HETBA1003866 27.257 12.805 22.440 12.069 15.414 19.103 9.229 7.5 HETBA1003868 95.701 54.991 58.923 31.090 41.733 59.461 48.174 43.4 HETBA1003879 62.950 44.572 159.217 48.098 42.446 37.097 36.010 45.8 HETBA1003880 134.462 70.074 103.271 50.699 47.956 67.668 44.498 30.5 HETBA1003880 134.462 70.074 103.271 50.699 47.956 67.668 44.498 30.5 HETBA1003885 77.675 69.096 172.368 55.129 49.424 41.309 24.247 31.5 HETBA1003885 76.675 69.096 172.368 55.129 49.424 41.309 24.247 31.5 HETBA1003893 38.6525 281.955 15.506 7.762 16.057 139.271 387.408 51.4 HETBA1003896 12.753 8.056 15.506 7.762 16.057 139.271 387.408 51.4 HETBA1003896 411.418 232.899 387.182 144.104 165.806 233.857 186.700 143.5 HETBA1003896 411.418 232.899 387.182 144.104 165.806 233.857 186.700 143.5 HETBA1003896 32.775 21.109 45.629 10.006 13.109 14.294 24.342 17.4 HETBA1003808 8.660 8.873 15.689 7.289 15.429 6.307 2.267 5.6 HETBA1003893 88.660 8.873 15.689 7.289 15.429 6.307 2.267 5.6 HETBA1003904 27.800 13.368 85.200.940 40.887 36.238 35.284 29.595 40.4 HETBA10039340 77.800 13.368 15.429 6.307 7.267 5.6 5.6 HETBA10039340 77.800 13.368 15.429 6.307 7.267 5.6 5.6 HETBA10039340 77.800 13.368 18.454 10.235 10.394 14.633 17.773 9.4 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 9.4 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 9.4 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 9.4 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 15.1 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 9.4 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 15.1 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 19.0 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 19.0 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 19.0 HETBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.773 19.0 HETBA1003950 73.300 13.000 14.000 73.000 18.80						8.662	11. 259	9.145	9.934
HETBA1003864   52.130   18.071   24.567   9.568   13.009   16.810   29.271   16.7   RETBA1003866   27.257   12.805   22.440   12.068   15.414   19.103   9.229   7.5   HETBA1003868   95.701   54.991   58.923   31.090   41.733   69.461   48.174   43.4   44.84								45, 837	48. 871
HEBBA1003866   27.257   12.805   22.440   12.069   15.414   19.103   9.229   7.5     HEBBA1003868   95.701   54.991   58.923   31.090   41.733   69.461   48.174   43.4     HEBBA1003860   134.462   70.074   103.271   50.699   47.955   67.668   44.498   30.5     HEBBA1003860   134.462   70.074   103.271   50.699   47.955   67.668   44.498   30.5     HEBBA1003864   99.190   48.465   73.499   34.796   54.399   57.269   63.551   68.8     HEBBA1003885   77.675   69.096   172.968   55.129   49.424   41.309   24.247   31.5     HEBBA1003890   12.753   8.056   15.506   7.762   16.057   139.271   387.408   5.1     HEBBA1003893   386.525   281.955   515.307   187.300   180.355   212.964   137.297   122.3     HEBBA1003896   411.418   232.899   382.182   144.104   165.806   233.857   186.700   143.5     HEBBA1003904   32.775   21.09   45.629   10.006   13.109   14.294   24.342   17.4     HEBBA1003908   8.660   8.873   15.689   7.298   15.429   6.307   2.267   5.0     HEBBA1003908   37.636   253.614   316.882   183.017   124.195   147.955   105.962   360.5     HEBBA1003939   28.064   25.844   35.675   20.306   20.378   19.070   16.457   15.689   16.293   10.006   13.109   14.294   24.342   17.4     HEBBA1003930   37.76   37.905   63.862   200.940   40.687   36.238   35.284   29.695   40.4     HEBBA1003930   37.765   37.907   3									16.795
HEMBA1003868 95.701 54.991 58.923 31.090 41.733 69.461 48.174 43.4 HEMBA1003879 62.950 44.572 159.217 48.098 42.446 37.097 36.010 45.8 HEMBA1003801 134.462 70.074 103.271 50.699 47.956 67.668 44.498 30.5 HEMBA1003803 99.190 48.465 73.499 34.796 54.399 57.269 63.551 68.8 HEMBA1003805 77.675 69.096 172.968 55.129 49.424 41.309 24.247 31.5 HEMBA1003807 60.203 22.185 33.582 16.896 21.181 29.281 31.275 22.8 HEMBA1003890 12.753 8.056 15.506 7.762 16.057 139.271 387.408 5.1 HEMBA1003896 411.418 232.899 382.182 144.104 165.806 233.857 186.700 143.5 HEMBA1003902 39.732 39.491 114.984 20.297 23.509 16.793 14.124 20.4 HEMBA1003904 32.775 21.109 45.629 10.006 13.109 14.294 24.342 17.4 HEMBA1003905 132.636 253.614 316.882 183.017 124.195 147.955 105.962 360.5 HEMBA1003937 87.005 63.862 200.940 40.687 36.238 35.284 29.695 40.4 HEMBA1003940 27.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HEMBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HEMBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HEMBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HEMBA1003940 77.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26.4 HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26.4 HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26.4 HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5.4 HEMBA1003950 8.31082 90.718 253.084 74.499 85.036 62.450 34.852 86.									7.524
HETBA1003879   62.950									43. 486
HEPBA1003880									45. 824
HEMBA1003884   99. 190   48. 465   73. 499   34. 796   54. 399   57. 269   63. 551   68. 8									
HETBA1003885									
HETBA1003897   60. 203   22. 185   33. 582   15. 896   21. 181   29. 281   31. 275   22. 8									
HETBA1003890									
HETBA1003893   386.525   281.955   515.307   187.300   180.355   212.964   137.297   122.3     HETBA1003896   411.418   232.899   382.182   144.104   165.806   233.857   186.700   143.5     HETBA1003902   39.732   39.491   114.984   20.297   23.509   16.793   14.124   20.4     HETBA1003904   32.775   21.109   45.629   10.006   13.109   14.294   24.342   17.4     HETBA1003908   8.660   8.873   15.689   7.298   15.429   6.307   2.267   5.5     HETBA1003926   132.636   253.614   316.882   183.017   124.195   147.955   105.962   360.5     HETBA1003937   87.005   63.862   200.940   40.687   36.238   35.284   29.695   40.4     HETBA1003939   28.064   25.844   35.675   20.306   20.378   19.070   16.457   15.5     HETBA1003940   27.800   13.368   18.045   10.235   10.394   14.633   17.733   9.5     HETBA1003941   57.997   16.835   24.582   17.381   15.884   23.428   19.757   13.5     HETBA1003942   38.168   19.747   45.852   32.660   22.333   24.695   10.791   21.5     HETBA1003950   8.366   8.726   5.814   3.195   4.756   3.396   8.814   5.5     HETBA1003950   8.366   8.726   5.814   3.195   4.756   3.396   8.814   5.5     HETBA1003950   8.366   8.726   5.814   3.195   4.756   3.396   8.814   5.5     HETBA1003950   53.133   29.785   31.879   18.932   16.178   21.708   32.094   35.									22.835
HEMBA1003896	HEPBA1003890				7.762				5. 124
HEMBA1003906	HEMBA1003893	386. 525	281.955	515.307	187.300				122.335
HEMBA1003902 39.732 39.491 114.984 20.297 23.509 16.793 14.124 20.4 HEMBA1003904 32.775 21.109 45.629 10.006 13.109 14.294 24.342 17.4 HEMBA1003908 8.660 8.873 15.689 7.298 15.429 6.307 2.267 5.6 HEMBA1003937 87.005 63.862 200.940 40.687 36.238 35.284 29.695 40.4 HEMBA1003939 28.064 25.844 35.675 20.306 20.378 19.070 16.457 15.4 HEMBA1003940 27.800 13.368 18.045 10.235 10.394 14.633 17.733 9.3 HEMBA1003941 57.997 16.835 24.582 17.381 15.884 23.428 19.757 13. HEMBA1003942 38.168 19.747 45.852 32.680 22.333 24.695 10.791 21. HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26. HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26. HEMBA1003958 12.870 13.019 20.678 7.159 38.521 442.120 272.494 21. HEMBA1003958 13.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003958 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003950 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.	HEMBA 1003896	411.418	232.899	382.182					143.577
HEMBA1003904   32.775   21.109   45.629   10.006   13.109   14.294   24.342   17.4     HEMBA1003908   8.660   8.873   15.689   7.298   15.429   6.307   2.267   5.6     HEMBA1003926   132.636   253.614   316.882   183.017   124.195   147.955   105.962   360.5     HEMBA1003937   87.005   63.862   200.940   40.687   36.238   35.284   29.695   40.687     HEMBA1003939   28.064   25.844   35.675   20.306   20.378   19.070   16.457   15.4     HEMBA1003940   27.800   13.368   18.045   10.235   10.394   14.633   17.733   9.5     HEMBA1003941   57.997   16.835   24.582   17.381   15.884   23.428   19.757   13.     HEMBA1003942   38.168   19.747   45.852   32.660   22.333   24.695   10.791   21.5     HEMBA1003945   59.457   32.900   46.079   23.037   21.163   36.632   32.279   26.5     HEMBA1003949   12.870   13.019   20.678   7.159   38.521   442.120   272.494   21.5     HEMBA1003950   8.366   8.726   5.814   3.195   4.756   3.396   8.814     HEMBA1003958   131.082   90.718   253.084   74.499   85.036   62.450   34.852   86.5     HEMBA1003958   13.1082   90.718   253.084   74.499   85.036   62.450   34.852   86.5     HEMBA1003959   12.105   11.228   18.520   6.548   7.960   18.122   12.612   10.5     HEMBA1003950   53.133   29.785   31.879   18.932   16.178   21.708   32.094   35.				114. 984	20.297				20.479
HEMBA1003908						13.109	14. 294	24.342	17, 444
HETBA1003936 132, 636 253, 614 316, 882 183, 017 124, 195 147, 955 105, 962 360, 962 105, 962						15.429		2.267	5.699
HETBA1003937 87.005 63.862 200.940 40.687 36.238 35.284 29.695 40.4 HETBA1003939 28.064 25.844 35.675 20.306 20.378 19.070 16.457 15.0 HETBA1003940 27.800 13.368 18.045 10.235 10.394 14.633 17.733 9.4 HETBA1003941 57.997 16.835 24.582 17.381 15.884 23.428 19.757 13. HETBA1003942 38.168 19.747 45.852 32.660 22.333 24.695 10.791 21. HETBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26. HETBA1003945 12.870 13.019 20.678 7.159 38.521 442.120 272.494 21. HETBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5. HETBA1003953 23.527 10.310 11.872 9.390 8.494 10.637 10.973 5. HETBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HETBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HETBA1003950 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.									360.995
HEMBA1003940 27.800 13.368 18.045 10.235 10.394 14.633 17.733 9.1 HEMBA1003941 57.997 16.835 24.582 17.381 15.884 23.428 19.757 13. HEMBA1003942 38.168 19.747 45.852 32.660 22.333 24.695 10.791 21. HEMBA1003945 59.457 32.900 46.079 23.037 21.163 36.632 32.279 26. HEMBA1003949 12.870 13.019 20.678 7.159 38.521 442.120 272.494 21. HEMBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5. HEMBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003950 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.									40.418
HEMBA1003940   27.800   13.368   18.045   10.235   10.394   14.633   17.733   9.4   14.633   17.733   9.4   14.633   17.733   9.4   14.633   17.733   9.4   14.633   17.733   9.4   14.633   17.733   1									15.626
HECBA1003941 57. 997 16. 835 24. 582 17. 381 15. 884 23. 428 19. 757 13. HECBA1003942 38. 168 19. 747 45. 852 32. 660 22. 333 24. 695 10. 791 21. HECBA1003945 59. 457 32. 900 46. 079 23. 037 21. 163 36. 632 32. 279 26. HECBA1003948 12. 870 13. 019 20. 678 7. 159 38. 521 442. 120 272. 494 21. HECBA1003950 8. 366 8. 726 5. 814 3. 195 4. 756 3. 396 8. 814 5. HECBA1003953 23. 527 10. 310 11. 872 9. 390 8. 494 10. 637 10. 973 5. HECBA1003958 131. 082 90. 718 253. 084 74. 499 85. 036 62. 450 34. 852 86. HECBA1003959 12. 105 11. 228 18. 520 6. 548 7. 960 18. 122 12. 612 10. HECBA1003950 53. 133 29. 785 31. 879 18. 932 16. 178 21. 708 32. 094 35.									9.868
HEMBA1003942   38.168   19.747   45.852   32.660   22.333   24.695   10.791   21.									13,795
HEMBA1003945   59,457   32,900   46,079   23,037   21,163   36,632   32,279   26,									21.900
HEBA1003949 12.870 13.019 20.678 7.159 38.521 442.120 272.494 21. HEBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5. HEBA1003953 23.527 10.310 11.872 9.390 8.494 10.637 10.973 5. HEBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEBBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEBBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.									26.903
HEPBA1003950 8.366 8.726 5.814 3.195 4.756 3.396 8.814 5. HEPBA1003953 23.527 10.310 11.872 9.390 8.494 10.637 10.973 5. HEPBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.								32.219	
HEMBA1003953 23.527 10.310 11.872 9.390 8.494 10.637 10.973 5. HEMBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.									21.625
HEMBA1003953 23.527 10.310 11.872 9.390 8.494 10.637 10.973 5. HEMBA1003958 131.082 90.718 253.084 74.499 85.036 62.450 34.852 86. HEMBA1003959 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.	HELBA1003950	8.366		5.814					5, 401
HEMBA1003958     131.082     90.718     253.084     74.499     85.036     62.450     34.852     86.       HEMBA1003959     12.105     11.228     18.520     6.548     7.960     18.122     12.612     10.       HEMBA1003960     53.133     29.785     31.879     18.932     16.178     21.708     32.094     35.					9. 390	8. 494			5. 252
HEMBA1003958 12.105 11.228 18.520 6.548 7.960 18.122 12.612 10. HEMBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.					74.499	85.036	62.450	34.852	86. 529
HEMBA1003960 53.133 29.785 31.879 18.932 16.178 21.708 32.094 35.							18. 122	1 12.612	10.591
11CCAN100300   33.100   23.100   01.013   101000									35, 333
									25, 303
									4. 985
[RELIDATION 3501   1.005   3.304   3.005   4.007   4.007   4.000   4.	LUESTIDA LUUSAB	1.039	1 3.300	3.304	2.033	1 4.034	1 7. 004	1.200	1

Table 28

	HEMBA1003968	40.219	26.894	55. 357	16.296	14. 511	28, 531	22.648	15. 420
	HEMBA1003974	147. 167	439, 547	139.030	117.010	33.973	54. 122	29. 356	338.820
	HEMBA1003976	20. 167	17.809	13.159	9.187	5,748	6.820	6.962	10. 367
5	HEMBA1003977	32.761	12.350	24.212	6.558	6.776	12.413	17.016	9.367
	HEMBA1003978	40.564	13. 858	10.812	11.585	11. 203	23.881	20. 489	17. 488
	HEMBA1003918	65, 803	34. 462	71.399	26.801	31.348	48.051	31. 355	42.728
			89.360	20.946	18.086	1.620	3. 781	3. 102	64. 356
	HEMBA1003982	15, 104		21.715	9.199	1.517	8.041	5. 977	7.559
	HEMBA1003985	15. 199	10.866					21. 302	24. 940
10	HEMBA1003987	48.695	30.080	108.473	25.632	23. 222	28.008	15. 392	
	HEMBA1003989	47.841	51.466	128.889	32.288	24. 298	24. 627 20. 604	16. 803	23. 174
	HELTBA 1004000	36.424	35.098	34.843	16. 292	19. 541		4. 943	9.742
	HEMBA1004006	8.411	42. 393	12.931	2.863	3.395	0.000		
	HEMBA1004007	135. 300	114.014	286.000	90.971	64. 473	74. 153	71.985	79.319
	HEMBA1004010	58. 331	152.845	38.786	18.676	18.819	35. 229	31. 514	80. 599
15	HEMBA1004011	62.306	16.294	38.336	12. 356	13.756	29. 583	26.091	7.986
	HEMBA1004012	47.010	38.053	139.110	42.415	22. 159	34. 340	27. 215	32.550
	HEMBA1004015	24. 416	26. 249	27. 372	12.243	13.962	25. 082	25. 133	12. 269
	HEMBA1004024	149.457	114.788	479.037	80.679	77.896	75.066	57. 366	93.859 19.376
	HEMBA1004029	81.485	31.944	43.520	19.897	20. 191	38.768	36.482	17.771
	HEMBA1004038	26.629	15.823	19.708	12. 109	7.832	14.400	12.855	10.405
20	HELIBA1004042	8, 177	10.678	12.830	6.612	11.484	7.963	11.320	17. 447
20	HEL'BA1004045	24.675	30.855	37.128	20.069	23. 538 40. 779	15. 509 45. 132	17. 299 47. 334	63. 844
	HEMBA1004048	95. 795	48.977	78.760	36.608 49.034	19. 297	56. 209	23. 320	68.865
	HEMBA1004049	55. 947	543. 954	47.428		25. 684		30. 423	32.553
	HEMBA1004051	69.776	31.608	51.948	13.046		38. 632	13.218	23.973
	HEL'BA1004053	29. 222	70.670	84. 481	24. 394	15.007	23.414	15. 237	
26	HEMBA1004055	39. 564	23. 202	34. 928	8.151	5. 353 81. 883	28.619 56.439	41.004	14.807 85.794
25	HEL/BA1004056	136. 121	122.072	413.353	75.363	4. 910	13.895	8.679	8. 388
	HEMBA1004060	17. 642 17. 144	11.826	29.995	9.507	8. 228	14. 145	12. 424	5. 810
	HEMBA1004067	165.029	79. 589	104. 390	62.419	50.783	89.115	94.004	91.850
	HEMBA1004071	28. 405	34. 722	37.707	19.775	14. 692	17.342	23.864	27.554
	HEMBA1004074	128. 445	51.388	148. 050	35.606	37, 851	50. 216	53.461	46.373
	HEMBA1004078	25. 126	14.714	20.940	9.721	16. 211	17. 398	17. 388	14.057
30	HEMBA1004085	42.006	24.067	36.862	15. 417	17.609	19.555	28. 362	21.993
	HEMBA1004085	27. 330	49.843	21.238	43.213	24. 232	16. 260	12.409	22.262
	HEMBA1004097	45.295	15. 292	27.795	13.971	26. 928	26.002	33. 192	19.361
	HEMBA1004100	40.930	37.210	48. 942	23. 245	10. 184	25.744	21.452	28. 594
	HEMBA1004103	101.036	101.281	184. 668	64, 176	44. 322	55. 385	41.050	40.000
	HEMBA1004110	89. 903	65. 107	57. 751	43.841	27.836	21.315	27.631	34.280
35	HEMBA1004111	171.907	134, 108	296.310	95. 474	115.874	78. 450	80.011	98.760
	HELBA1004124	177. 408	71.838	103.065	37.865	46.198	68. 531	109.364	77.083
	HEMBA1004130	64. 543	54. 797	171.602	50.628	35. 382	25. 601	19.599	23.097
	HEMBA1004131	41.654	24. 184	33. 975	26.913	23.365	28. 790	20.022	24. 999
	HEMBA1004132	55. 906	42.840	162. 243	42.708	30, 251	28.863	19.780	22. 237
	HEMBA1004133	64. 624	30.838	38. 522	29.390	20.897	28.027	28.747	33. 333
40	HEMBA1004138	61.197	21.853	23. 858	17.376	9.337	30.080	17.345	22.082
	HEL18A1004143	15.715	9.656	21. 209	10.565	10.539	14.067	11.441	9. 994
	HEMBA1004146	40.893	21.789	90. 537	30.633	32.870	23.542	14.368	20.982
	HEL'BA1004148	59.990	18.795	22. 167	11.049	17.531	18.309	29.374	22. 628
	HEL'BA1004149	16.284	11.131	18.385	7.758	7.634	7.677	5.890	13.683
	HEMBA1004150	5. 223	4. 403	4. 468	3.044	2.553	2.158	2.052	2. 260
45	HEMBA1004154	111.110	40.836	69.965	31.437	46. 253	58. 472	62.983	47.866
	HEMBA1004164	139.670	107.565	315. 189	77.326	47.327	57.372	46.726	67. 257
	HEMBA1004168	24. 042	16.530	18. 698	9.347	9.400	13.838	3.054	13.060
	HEDBA1004199	22.894	9.047	10.461	8.631	7.704	7.849	6.889	7. 253
	HEMBA1004200	33. 301	51.362	83.462	26.185	27.548	17.580	17.235	32.109
	HEMBA1004201	54.766	23.783	32.370	17.449	21.835	22.123	25.993	20.006
50	HEMBA 1004202	14. 526	10.484	12.784	6.804	5.704	9.594	8.672	11.673
	HEMBA1004203	47.655	20.140	34. 882	13.604	14.171	19.946	16.079	18.151
	HEMBA1004207	6.344	3. 206	11.421	3.936	6.145	5.704	21.692	7.780
	HEMBA1004210	33.071	43.543	33.120	16.340	41.396	21.814	19.639	15.015
	HEMBA1004225	73, 182	63.749	226. 133	59.565	43.156	32.703	25. 781	40.078
	HEMBA1004227	83.820	31. 222	42.541	16.931	17.785	28. 177	25. 468	30.978
55									

#### Table 29

					010 2)				
	HEMBA1004235	99.954	57, 144	62.536	27.672	34.345	69.613	47, 182	38.807
	HEMBA1004237	27.504	21.542	17.029	18. 289	11.697	19.212	12.031	16.922
,	HEMBA1004238	79.210	38. 454	102.493	34, 130	27.841	36.089	27.438	34. 578
5	HEMBA1004241	5.663	2.654	7, 035	2.556	1.072	2.912	4. 422	1.294
	HEMBA1004242	256.862	65. 757	191.327	80.010	76. 455	85, 478	89.242	62.567
	HEMBA1004243	72.699	55. 276	60.764	28. 287	47, 148	36, 800	28, 491	47.743
	HEMBA1004246	44. 915	30. 967	100, 300	22,414	17. 109	15, 470	12.686	18.700
		66.750	16. 238	24.674	18. 889	22.763	31.897	38.415	17.377
	HEMBA1004247	13. 953		17. 581	11.953	11.378	14. 538	12.794	9. 562
10	HEMBA 1004248		18. 412		6.401	5. 142	14. 218	12.652	11.966
	HEMBA1004250	24. 439	10. 494	10.631		9. 550	14, 570	21.841	18. 200
	HEMBA1004252	37. 349	20.650	22. 246	9, 949		13. 845	11.265	19. 838
	HEMBA1004260	10.994	19. 320	16.415	15.707	20.374	12. 282	15.089	11.397
	HEMBA1004264	22.716	14.715	13.358	7.615	5. 234	115. 073	102.973	144, 125
	HEMBA1004267	235.310	195. 750	654. 331	171.071	174. 292			
15	HEHBA1004272	28.776	19. 025	23, 678	13.063	12.012	15, 529	14. 123	14. 593
	HELBA1004274	62.157	50, 491	53, 598	30. 356	36. 472	42.005	58.020	51.617
	HEMBA1004275	70. 423	38. 514	45. 176	17.443	18. 132	34. 031	36.295	22. 171
	HEL:8A1004276	33.630	4. 481	14.011	9. 548	9.099	14. 035	10.406	8.615
	HEMBA1004279	16.536	11.082	13.356	14.834	7. 333	10. 255	8.919	12.068
	HEMBA1004284	29.688	30. 297	64.483	13.658	17.646	17. 327	17.630	13. 770
20	HETBA1004286	32. 471	16.566	18.049	12.391	6.773	17. 625	23.811	13. 547
20	HEMBA1004289	81.573	62.930	165.571	49.704	34.785	37.379	28. 939	41.740
	HEMBA1004293	72.466	34, 902	48, 669	32.705	17.408	57.764	53.695	45.065
	HEMBA 1004295	37. 595	12. 116	29.975	11.634	5.514	25.018	23.797	20. 926
	HEMBA1004302	10.880	5. 912	7.885	10.025	5. 190	6.060	5. 264	9. 355
	HEMBA1004306	426.811	177.321	335. 168	107.646	123. 947	256. 397	251.772	134. 005
	HEMBA1004312	37.953	30.864	105. 533	30.747	25.847	16.140	16. 283	24. 272
25	HEMBA1004314	29. 396	23. 332	95. 584	22.179	18.544	11.015	8.804	20.974
	HEMBA1004321	47.670	29, 150	105, 316	35.655	23.139	31.309	29.736	47.858
	HEMBA1004323	87.295	65. 931	221.440	44.690	41.425	36.609	34.117	39. 135
	HEEBA1004327	65.869	21. 284	21.540	11.985	14.419	27. 213	27.030	20.118
	HEMBA1004329	67.920	44.687	132.755	12.977	21.556	32.356	17.478	26.773
•	HE™BA1004330	8.765	7.655	16.827	7.164	3.843	9, 511	7.660	4. 615
30	HEMBA1004334	16. 438	21.355	31.680	15. 109	26.670	13.368	10.581	13. 568
	HEEBA1004335	204. 961	102.859	325.226	69. 979	64. 392	78.772	71.641	83.525
	HEE8A1004341	186.677	30. 208	61.439	15. 995	41.404	87.221	89.558	40. 224
	HEI:BA1004344	261.576	76.316	123.332	42.705	51.432	26.797	42.054	59.071
	HEMMA1004347	65. 249	32.610	97.858	37.038	21.953	33.115	33.526	36.846
	HEEBA1004349	22.353	35.727	29.441	19.803	18.786	23.126	19, 103	18.719
35	HEMBA1004352	75.508	65. 544	237.050	49.039	34, 141	32.597	28.166	46.343
	HEMBA1004353	54.322	66.042	132.169	40.563	27.380	39.551	30.556	56.886
	HEEBA1004354	43.687	29. 352	79. 264	22.784	20.533	21.755	16.860	22.429
	HEMBA1004356	44.730	22. 201	27.487	10.404	8.280	22. 159	16.039	15.038
	HEMBA1004360	91.412	28. 429	71.634	26.232	36. 259	59.602	38.361	50.410
	HEMBA1004366	9.956	10.099	14. 263	5.481	5. 631	6.802	6.791	6.167
40	HEMBA1004372	3,613	4. 593	5. 338	0.000	1.638	1.507	3.555	1.568
40	HEMBA1004377	53.834	41.410	47.048	29.140	26. 163	34. 545	30.827	33.572
	HEMBA1004389	20.540	22.800	24.474	14.497	13.968	16.620	14.951	17.114
	HEM8A1004391	60.284	22.653	44.013	14.283	19.018	31.716	23.931	23.617
	HEMBA1004393	177.786	197.548	108, 554	32.455	75. 399	76.587	39.777	44.665
	HEMBA 1004394	28. 949	11.849	12.442	5.544	10.440		10.981	8.836
	HEMBA1004396	37.907	26. 956	102.760	18.571	16.519	15.025	13.681	21.980
45	HEMBA1004401	22.519	21.858	30.601	14. 945	13.592	15.418	20.530	20.774
	HEMBA 1004405	42.933	38.835	117.844	34.528	23.557	19.155	18.506	29.842
	HEMBA1004408	50, 497	27, 151	55,000	25.559	15.351	19.522	15.546	20.863
	HEMBA1004414	45.769	51.722	64.316	19, 655	19.324	39.735	26. 527	36.385
	HEMBA 1004429	61.867	59.067	190.058	39.014	50. 304	38.462	27.517	46.317
	HEMBA 1004433	49. 568	39. 828	146. 938	37.521	28. 383	24. 241	24.651	42.005
50	HEMBA 1004440	31.849	22, 499	37. 132	18.742	22.366	23.183	21.969	35.073
	HEMBA 1004444	59.488	46. 586	163.763	10.695	29.990	24.833	22.908	37.635
	HELBA 1004446	22.134	12.309	29. 426	11.920	3. 385	14.862	10.855	16.078
	HEMBA1004451		21.261	28. 136	18. 194	15.678	19.695	21.159	
	HELBA 1004452	31.688	5. 268	18. 479		5. 737	17.680	14, 173	
	HEL'8A1004454		27. 897	32.786		20. 631	24. 494	22.897	29.042
55	UCT-041004434	50.056	1 41.031	1 32.100	1 24. 302	1 20.001	1 5 41 767		
55									

Table 30

	HEMBA1004460	138,550	96.143	356.058	74.883	54.735	70.698	38. 344	55.945
	HEMBA1004461	64.074	19. 163	24. 808	12.846	16.373	23, 508	22.827	10, 137
	HEMBA1004468	134.439	72.774	210.409	77.409	60.142	56.229	42.361	49. 457
5	HEBBA1004479	82.994	32.899	92. 282	35.519	44. 435	43, 183	31.836	37.116
	HELBA1004482	5, 602	7.682	11.248	36.034	2.926	5. 535	5.693	5.972
	HEMBA1004491	16.736	6. 285	17.615	14.018	11.729	19.804	10.683	11.280
	HEMBA1004499	94, 095	71, 456	148. 355	58.479	48.596	46.968	46.648	57.279
	HEMBA1004502	21.523	9. 344	18.265	7.282	9.979	2.762	3. 174	13. 389
	HEMBA1004505	26.042	15. 980	43.855	17.516	15. 469	22.190	22.873	15.812
10	HEMBA1004505	12.004	29. 395	42.664	21.849	21.426	14.469	9. 224	9.845
	HEMBA1004507	96.377	87.688	99. 177	103. 472	34. 160	81.068	54, 939	151.142
	HEMBA1004509	52.657	14. 880	19.120	16. 228	17.009	24. 783	24. 565	13.476
	HEMBA1004523	20. 156	18, 209	11.197	16.529	14, 651	13.004	20. 267	19.467
	HEMBA1004528	42.620	27.819	48.069	14. 426	25. 267	43.038	40.239	37.718
46	HEMBA1004534	75.090	41. 159	44. 399	31.300	16.686	31.317	21.009	18.589
15	HEMBA1004536	31.531	13.343	23. 564	14.085	4.408	13.133	12.981	15. 201
	HEUBA1004538	352.363	181.508	233.819	97.018	122. 402	183.507	100.197	150.062
	HEMBA1004542	47. 360	17.733	29. 238	17.280	12.324 39.674	17. 317 38. 686	22.764 23.830	15. 212 33. 542
	HEYBA1004552	63. 401	29. 585,	26.857 25.084	43. 567 16. 273	32.820	33.809	20.812	5. 536
	HELBA1004554 HELBA1004558	62. 231	11.953 12.133	31.036	15.840	30.638	65. 183	19.155	30. 921
20	HEMBA1004560	30.217 68,901	16.566	17. 908	16.431	10.034	30.093	22.379	22. 583
	HEMBA1004564	48, 119	14. 911	35. 565	31.983	32.464	30.028	20.965	32.479
	HEMBA1004566	32, 479	29. 553	20.970	32.788	42.949	40.715	23. 273	32.960
	HEMBA1004573	17, 728	13, 843	7, 118	9.972	19.952	9.755	9. 278	8.100
	HEMBA1004576	39. 572	26.733	42.044	10.704	37.505	32.441	17.232	25.361
	HEMBA1004577	46.233	11.570	97.881	39.434	13. 437	41.089	34. 426	35.314
25	HEMBA1004586	82.532	71.398	213.814	70. 289	52.589	45.729	23.395	38.312
	HEMBA1004596	72. 534	32.493	45. 820	27.585	27.854	34.997	33.847	38.473
	HEMBA1004604	99.019	48. 582	103. 587	36.723	49. 392	48.377	56.558	69.256
	HEMBA1004607	53. 557	37.013	100. 999	27.559	26.143	28.796	21.692 8.858	42.044
	HEMBA1004610	20.690	14.854	69. 908	15.349	12. 120	9, 108	6.697	15.087
	HELBA1004617	22.592	20. 386	42. 426 209. 059	22.819 49.931	15. 568 29, 836	10.691	12.194	27. 438
30	HELBA1004622	78.025	46.803 36.312	110.684	22.791	14.118	17. 193	15. 579	20.821
	HEMBA 1004626 HEMBA 1004629	38. 170 33. 858	37.886	87.440	53. 228	47.341	28. 160	12.170	28.096
	HEPBA1004631	35. 946	10.475	4. 434	7.390	17. 128	22.775	9. 569	32.852
	HEMBA1004632	27.084	13.891	23. 598	10.209	7.802	11.754	22.566	6.362
	HEMBA1004633	78. 391	33.135	114.054	17. 197	49.008	60.659	48. 857	40.810
25	HEMBA1004536	52. 397	20.706	34.962	10.085	22.609	21.255	13.502	25.039
35	HEMBA1004637	4. 228	4.304	6.747	5. 278	9.756	4.086	2.597	5.024
	HELEBA 1004638	0.241	0.000	0.000	1.008	0.000	0.000	0.113	0.000
	HEMBA 1004645	57. 971	29.263	111.067	32. 645	17.998	27.214	20.560	24.845
	HEMBA1004656	16. 139	9.194	21.399	12.766	18. 216	14.099	17. 122	12.004
	HEMBA 1004657	20. 820	23.742	69.842	9.422	138.932	42.697	9.048	13.383 7.512
40	HEMBA 1 004666	7. 321	3. 174	18.097	5.962 30.591	9.830	28.018	25. 500	25. 624
,,,	HEMBA 1004669	94. 910 57. 231	36. 291 17. 070	60.538	23. 280	13. 173	24.312	23.413	14. 342
	HEMBA1004672	63. 471	50.154	146.619	39.883	31.559	25.617	20.328	28.099
	HEMBA1004689	152. 993	93. 435	103.311	81.212	50.901	83.998	57. 329	84.276
	HEMBA1004690	28.240	10.247	13.401	8.159	4. 952	13.963	13.991	11.785
	HEY BA1004693	18.359	15. 228	20.803	14. 290	13.070	16.726	9.014	13.531
45	HEMBA1004897	81.532	48.847	148. 587	58.849	34.416	51.983	42.641	50. 271
	HEMBA1004702	97.518	62.966	49.904	20.714	42.224	58. 936	64, 906	37.506
	HEMBA1004704	99.561	48.717	236. 687	38.866	33.457	38.377	24. 626	31.783
	HELMA1004705	12.717	12.313	40.950	9.649	17. 803	10.638	5. 969	4.810
	HEMBA1004706	33.616	9. 825	16.175	10.779	10.830	17. 906	13.036	12.703
	HEP3A1004709	51. 126	39. 934	136.723	32. 285	25.072	21.674	15. 230	23.755 9.790
50	HEMBA1004711	46.766	9. 203	57.020		14. 304	16.154 65.719		52,009
	HEMBA1004723	121. 283	47.643	73.497	30.236	56.917 48.208	34.021	35. 739	12.501
	HEMBA1004725	56. 905 36. 072	32.051 10.037	30.016	7.633	13.361	7.545		34. 832
	HEMBA1004730	30.769	29. 884			2. 998	8.055		2.822
	HELBA1004734	11. 912	11.974			12. 556	7. 653		
E E	TILLADA 10041 34	1 11.312				•			
55									

Table 31

					72 653 1	40 000 1	02 516 1	10.070	24 600
	HEMBA1004736	55. 309	25. 331	132, 333	45.653	38.696	23.516	19.970	34. 509
	HEMBA1004748	53.832	20.004	172.069	29. 562	22. 161	14.904	12.665	18.349
5	HEMBA1004749	127. 285	45. 137	73.698	27.788	33. 184	60.214	44.636	42.250
•	HEMBA1004751	81.283	64.830	173.888	54, 165	36.368	41.802	31.283	43.505
	HEEBA1004752	59.058	32.785	109. 428	32.254	29.090	34. 259	30.970	33.029
	HEMBA1004753	204.044	247.466	406, 165	156.689	102.755	131.323	83.515	263.882
			59.677	83.850	22. 148	29.800	30.642	13.064	23. 261
	HEMBA1004755	57.838					9. 109	111.628	14. 885
	HEMBA1004756	9.965	16.228	11.023	8. 349	6. 780			
10	HEMBA1004758	36, 487	26.558	116.970	22. 341	14.553	14.773	11.840	14.406
	HELBA1004763	67.343	19.641	33.742	13.841	16.720	25. 489	23.061	18.650
	HEMBA1004768	29. 177	24.043	38. 303	6.673	10.298	3. 197	10. 352	13. 391
	HEMBA1004770	10.327	14. 492	10. 901	6.416	6.310	7.963	10.868	7. 955
	HEMBA1004771	46.910	34.314	76.491	31.609	22.830	23. 102	30.433	32. 358
	HEMBA1004775	39. 253	28, 706	63.968	24. 931	18.754	43.049	32.720	26.795
	HEMBA1004776	22.604	11.017	10.103	5.466	9,000	16.400	10.105	8. 046
15	HEMBA1004778	78.144	77.681	223, 475	37.540	33, 791	32.337	24.067	43. 529
			18. 370	102.812	8. 313	15. 151	11.373	9.479	6. 329
	HEMBA1004784	9.826					10.674	13.732	11.615
	HEMBA1004785	25, 723	16. 345	26.216	6.651	10.649			
	HEMBA1004789	18. 173	14. 5081	16.096	7.804	8.691	10.011	7.713	11.389
	HEMBA1004795	14. 283	12.973	25. 122	11.028	9. 351	9.757	9.905	12.028
20	HELBA1004797	65. 927	33.745	73.888	34.142	28. 246	40.067	32.715	25. 583
20	HEMBA1004803	36.634	41.124	65.880	27.072	30.957	22.607	22.520	26.554
	HELBA1004805	11.997	8.183	21.467	8.868	9.653	9.000	7.894	8: 399
	HELBA1004807	16. 352	14.481	22.459	11.249	12.009	13.340	7.935	9.118
	HEMBA1004816	29.782	24.075	95.884	18.110	29.259	8. 180	12.578	10. 934
	HEL BA1004820	8.636	7.466	8.862	4. 249	4.018	4, 269	6.876	3.493
	HEMBA1004833	159. 947	50.729	81.248	38.650	64.754	83. 155	56.657	65, 121
25	HEL'BA1004847	51.456	25. 570	40.694	21.115	36.572	35. 053	31.537	40. 529
25		77.254	24.014	38.620	21.854	26. 080	54. 413	50.197	24. 185
	HELBA1004850	11.234		72.480	23.951	31.887	25. 058	20.050	20. 982
	HEMBA1004863	57, 117	32.704					20.308	17.843
	HEYBA1004864	46.043	27.344	59, 824	26.750	13.898	16.719		
	HEMBA1004865	12. 257	14.642	31.748	44.090	14. 331	13.454	13.835	15.797
	HELBA1004880	56.788	50.021	126.837	35. 420	26. 589	24.064	20.647	23, 264
30	HEMBA1004882	42.450	18, 453	29. 340	16.782	13.013	13.652	10.676	19.977
	HEMBA1004885	8. 545	4. 947	5.350	4. 891	2. 933	3.711	3.652	6.615
	HEMBA1004889	28.103	22.485	32.049	17.078	14. 363	23.391	15.605	16.916
	HEMBA1004900	19.922	15.709	33.254	10, 423	9.045	6. 539	5. 245	9.440
	HEMBA1004909	88. 522	49.269	163.284	48. 147	35.537	36.045	18.861	27.933
	HELBA1004918	64.384	43. 134	105.868	34.899	22.323	24.073	15.857	25. 370
		47.731	37.996	69.168	19.659	26. 441	18. 192	10.213	20.111
35	HELBA1004923		14.003	10.808	12.050	7.539	9. 882	8.967	11.809
	HELBA1004929	11.048		279.652		66.618	34. 331	31.091	41.874
	HEMBA1004930	101.277	92.425		80.664			96.467	5. 417
	HEMBA1004933	9. 145	5. 566	12.895	7.786	12. 296	10.327		12.805
	HEL'BA1004934	7.311	7. 106	43.966	10.208	4.750	5.866	9, 143	
	HEEBA1004937	43. 331	27.219	38.802	15. 368	17.734	15. 280	15.784	46. 365
40	HEMBA1004943	51.072	26.833	32.001	21.614	16. 458	27.585	29.628	38. 533
40	HEMBA1004944	84. 363	46.788	126. 294	43.803	28. 989	38. 514	31.589	23.074
	HEMBA1004946	64.638	28. 144	37.908	17.163	24. 332	27.854	34.636	31.712
	HEMBA1004952	90.835	18.893	40.862	12.824	20.090	33.568	20.052	19.020
	HEMBA1004954	14. 656	36,003	41.485	27.126	23.696	20.777	6.945	29. 261
	HEMBA1004956	5.975	9. 923	6.635	7.743	0,953	4.578	1.565	5.188
	12412 1 4 2 4 1 4 4		77.420	136.061	60.735	49. 221	47.560	29.646	45. 929
45	HECBA1004950	31.046	5. 439	7.559	12.468	17.946	16.068	19.705	18. 480
	HELBA1004972	77.318			35.819	27. 295	40. 233	30.004	50.710
	HELIBA1004973		13.502	16.731	9.641	11.726	14.716	19. 197	22, 580
		35. 524						2.217	10.523
	HELBA1004977	6.756	9.870	11.419		29.373	8.701		
	HEMBA1004978	8.689		13.909		5. 158	5.699	2.642	10.106
	HEMBA1004980	34.093		87. 268	25. 974	18.071	16.453	11.605	22. 124
50	HEYBA1004982	14.750		17. 944		8. 250	11.553	6.083	
	HEL'8A1004983	38.285	13. 488	20.831	11.831	3.348	10.309	11.455	
	HEYBA1004995	27. 256	28.515	26.297	18.434	25.474	22.491	24.452	33.683
	HEMBA1005004	13.855		33. 238		7.816	13.134	7.576	14.698
	HEMBA1005008	64.714						18. 581	
	HEMBA1005009	34. 543							
	LITTION LOCOTON	34. 343	13.013	1 , 3, 702	1 . 5. 573	1 11 16 6			

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Table 32

	IUPMPA16ACATA	40 350	04 072	25 240	20 924	20 144	22 620 1	20.777	27.016
	HEMBA1005019	49. 260	24. 872	25. 349	20.834	30.144	32.629		
	HEMBA1005021	37.224	63.713	38.065	17.061	15.647	20. 369	25. 859	32.656
5	HEMBA1005029	30.265	17.783	35. 352	16.531	19.588	26.517	15. 798	16.604
•	HEMBA1005035	393. 404	200.167	574. 746	133.872	210.689	177.872	156.563	170.510
	HEMBA1005035	115. 345	41.961	73.015	39. 541	44. 451	66.623	55.833	51.349
	HEMBA1005039	28.850	19. 922	57.018	13.971	11.999	26. 427	15. 206	7.350
	HEMBA1005047	93.995	31.868	54. 335	18.576	28. 338	31.562	31.930	23.751
				73. 330	29.830	26.504	35. 887	21.640	35.653
	HEMBA1005050	78.015	41. 590						
10	HEMBA1005062	23.050	15.803	29.553	15.707	7.836	15.618	19. 435	13. 336
	HELBA1005066	10.980	11.364	31.553	13.509	5.668	10. 541	5.005	10.849
	HEMBA1005067	39.308	34. 578	39.795	44.519	24.643	21. 272	19.379	20.121
	HEMBA1005070	73.155	34. 949	68.556	29.956	38.004	38. 211	48.007	31.733
	HEMBA1005075	88.089	37.798	148.675	40,537	33.271	33.074	28.661	30. 201
	HEMBA1005078	100.064	37.746	66,827	33, 115	41.170	55, 560	51. 231	17. 112
	HEMBA 1005079	137, 757	86. 238	294, 118	73.304	76.035	75.084	47. 255	76.170
15	HEMBA1005083	18. 102	7.642	17.087	6.711	6.184	8. 675	9. 287	9.609
	HEMBA1005084	82.712	38. 248	47.063	26.664	27. 435	37, 552	38. 419	28. 349
						18.926	23.875	8.895	12.447
	HEMBA1005088	31,610	22. 435	76.774	22.700				
	HEMBA1005089	68. 944	55. 156	178. 226	34.742	32.350	38. 645	22.869	28. 148
	HEMBA1005090	148.861	86.156	117.997	94.811	57.034	81.098	54. 187	116.066
	HELBA1005096	83.125	30, 911	63. 940	33.378	33.962	48. 589	35, 467	36.021
20	HEMBA1005101	69.080	14.020	34. 136	10.197	13.998	34. 420	22.696	15.975
	HEMBA1005107	82.659	25. 203	36.223	11.215	21.514	32.720	25. 972	21.337
	HEMBA1005113	7.977	17. 225	31.501	7.563	44.493	5. 157	6.957	9.761
	HEMBA1005123	173.637	77, 260	555.672	126, 908	94,628	90. 446	70.735	90.016
	HEMBA1005133	58. 192	40. 749	122.920	29.864	16.700	17.652	8.802	18.988
	HEMBA1005135	8. 259	9. 125	14. 962	2.213	16.732	6.892	3.383	6.189
26	HEMBA1005145	185. 299	101.220	352.159	92.082	88.750	122.118	76.475	90.044
25					120.663	125. 192	96.704	92.083	128.030
	HEMBA1005149	220. 122	109. 352	274.492					
	HEMBA1005152	125. 948	96. 291	226.882	58.505	33.738	46. 323	27. 534	34. 457
	HEL:8A1005159	15.760	11.274	9.399	6.198	6. 191	6.861	12.001	4.556
	HEYBA1005172	1653. 208	89.658	73.666	54.667	33.118	55.680	32. 520	70. 907
	HEM8A1005185	9. 954	17, 248	10.492	34.452	3. 558	3. 117	6.026	11. 173
30	HEYBA1005186	23.745	10.048	27.091	13.067	7,719	15. 412	15.086	15. 591
50	HEY'8A1005195	14. 573	8.648	11.038	19.306	6.313	25. 313	13.510	9. 183
	HEMBA1005201	52. 322	13. 197	47.505	13.091	12.078	8. 531	23.532	9.848
	HEMBA1005202	98. 566	30.141	71.588	27.954	44, 381	66. 294	42.390	39.695
	HEMBA1005204	184. 429	287.156	382.039	168.753	203.458	222.970	143.609	358.646
	HEMBATO05206	148. 946	61.309	84. 791	34.139	49.115	66. 295	83.608	76, 159
	HEMBA1005219			9.606	8. 236	8. 038	13. 031	7.751	11.441
35		21.685	17.755				53.017	26.025	32.004
	HEMBA1005223	79.969	42.665	113.460	40. 547	32.099			
	HEYBA1005229	26.819	9. 926	21.841	3.135	5, 090	6.656	4.681	7.079
	HEMBA1005230	71.184	67.313	201.065	79. 279	59.679	77. 484	47.808	66.511
	HEL:3A1005232	7.374	6.386	17.522	8. 552	3. 285	12.098	4. 975	3.965
	HEPBA1005238	96.780	44. 134	51.932	8. 128	20.776	69. 291	49.474	35.019
	HEYBA1005241	142.598	104. 185	428.635	78.773	78.033	74.434	42.333	63.097
40	HEP301005244	75.771	32.597	37.797	16.459	12. 489	35. 934	31.814	35. 602
	HEMBA1005246	241.316	60.348	73.077	25.067	41.351	117.666	88. 193	54.014
	HEMBA1005251	37.505	33. 247	108, 631	23.585	14.915	23. 393	14.302	16.409
	HEMBA1005252	53.401	25. 532	37.199	15, 002	20.744	31.279	24.207	27.562
	HEL-BA1005267	17. 238	39. 564	20.097	27. 506	31.874	11.013	14, 526	14.024
	THE SECRETARY AND ADDRESS OF THE PARTY OF TH		·			10 100	11 045	8.914	11. 978
	HEPBA1005274	16.538	8.744	18.308	9.021	10.103	11.943	25.040	
45	HEPBA1005275	69.133	43. 329	216.468	46. 290	57.647	37.411		41.913
	HEMBA1005288	65.401	50. 495	150.714	33.833	34.633	28. 241	24, 910	40.164
	HEMBA1005293	17. 403	9. 430	23. 201	4.467	3. 192	25. 620	6.775	8.771
	HEMBA1005296	223.097	811.623	894.835	738. 361	220. 523	698. 319	418. 435	1376. 785
	REMBA1005301	36.708	16.970	29.798	11.929	12.544	22. 221	35.726	32.270
	HEMBA1005304	83.978	71.914	260.016	50.686	36, 101	36. 160	24.896	47.838
50	HEMBA1005305	44.218	33.773	74.215	27.494	27. 352	34. 920	21.424	38.882
	HEMBA1005311		20. 140	48. 263	13.836	7. 908	8. 958	9. 090	10.440
	HEL:BA1005313	33.034		17.550		11.502	14. 209	63.072	9. 124
		11.165	36.175		7.047				
	HEPBA1005314	6.948	2, 955	22.604	6. 336	3.812	5. 787	4.069	4.891
	HEL 341005315	72.349	54, 139	156.842	34.545	43.132	26.415	28. 942	28.442
	HELBA1005317	20.230	8. 451	17.258	9. 796	11.664	9.976	9.263	8.017

Table 33

								4 046
HEMBA1005318	14.755	5. 931	13.883	5. 228	5.376	9.013	5. 511	4.845 17.915
HEMBA1005324	98.070	33.348	44. 270	26.052	35.446	48. 523	30. 889 16. 942	29.580
HEMBA1005331	24. 826	335. 211	15.947	26.496	14.744 20.830	21.427	32. 481	44. 585
HEMBA1005337	19.080	18.022	19, 429	5. 217			40. 921	36. 285
HEMBA1005338	61.533	38.788	63.113	23.657	30. 437 72. 524	48. 455 167. 620	135. 992	68. 042
	384. 481	88. 937	143.574	53. 983	53. 484	67.048	30, 456	42.612
	111.629	68.949	220. 401	62.090		34. 233	21.666	47. 596
HEMBA1005359	87.635	64. 332	175. 543	59.707	36.743 21.917	19. 092	20.883	12.720
HEMBA1005362	25.674	25. 093	18.642	30.797	19. 753	5. 180	2.877	7. 198
HEHBA1005364	6,677	2.817	5. 168	13.116	30, 138	27. 987	16, 766	22.415
HEMBA1005367	51.911	28. 536	74. 559	28.446	9. 177	6. 402	9, 312	4. 913
HEHBA1005372	11. 289	6.819	11.700	5.659	30. 987	24. 792	23.695	30.728
HEMBA1005374	64.639	57.505	120, 218	32.738 8.862	7,648	11. 978	9. 051	4.019
HEMBA1005379	29. 549	13.813	12.040	70.213	26. 226	53, 452	88. 235	85. 480
HEMBA1005382	140.116	94. 743	104.609	10.250	8, 543	11, 030	7. 498	9.010
HEMBA1005384	33.109	15. 221	21.713	29.541	31.691	44. 619	35. 179	29. 136
HEMBA1005386	111.062	30. 547	52.790	42.528	35.894	16.765	11.513	24. 601
HEMBA1005389	66.821	32.429	129. 272	25. 833	25.639	30. 857	16, 944	24. 443
HEHBA1005394	35. 794	18. 327,	22.715	14.621	15. 504	27. 461	15, 586	32.390
HEMBA1005403	40. 404	14.030	54.041	44. 257	67.383	35.010	23. 590	44.612
HEMBA1005408	51.701	45.069	71.813	12.035	10.197	6. 188	3. 955	8.910
HEMBA1005410	4. 534	4. 269	163,001	67, 133	50. 499	41.243	22.652	35, 008
HEMBA1005411	75. 220	94.039 26.430	69. 138	35.773	15. 442	19. 286	14. 057	23, 010
HEMBA1005423	35.745	12.073	14, 418	5.345	11.591	8. 954	3.082	7.203
HEMBA1005426	14. 366	99. 596	61.088	47.865	59. 821	53.861	25. 223	46.397
HEMBA1005430	66. 444 52. 945	15.385	36. 316	19. 210	23. 854	37.895	19.556	18. 127
HEMBA1005438	51.806	28, 359	33. 314	17.787	19. 295	21.754	13.422	29. 941
HEMBA1005443	108. 954	165.667	426. 408	91.550	77.559	76.024	105.042	108. 232
HEMBA 1005447	51. 383	39, 578	65. 244	29, 171	28,000	21.457	18,763	23.755
HEMBA 1005449	86. 452	20. 253	41. 861	15, 939	27.647	39.311	28.567	27.508
HE BA1005452	110.567	52.128	74.119	42.532	39.847	53.326	67. 529	72.233
HEMBA 1005454	7.997	16.821	17. 998	14. 293	14.436	8.454	6. 498	11.445
HEMBA1005468	185.066	78.008	126.372	56.026	56. 490	78. 922	61.083	57.511
HEMBA1005469	88.419	54, 761	196.280	63.682	53.661	42.639	23.441	30, 144
HEMBA 1005472	37. 878	41.710	88.807	34.196	28. 126	21.983	24. 350	30. 575
HEMBA1005474	89. 169	55.263	212.086	51.664	50.480	66.508	39. 590	30.322
HEMBA1005475	212. 273	98. 359	182.707	110.945	105. 968	98.316	56.095	68.647
HEMBA 1005489	61.603	40.439	42.459	21.361	21.335	31.130	11.578	25.898
HEMBA1005497	10.325	12.396	5. 705	8.252	5.611	9.949	2.213	16.039
HEMBA1005500	86.636	39.755	180.843	46.031	28.654	31.809	14.951	31.189
HEMBA 1005506	24.029	3.468	17.794	7.400	5. 464	9.419	6.593	3.51
HEMBA 1005508	12. 944	12.524	22. 247	8.536	16.857	11.561	7.741	12.05
HEMBA 1005511	116. 338	59.193	267.636	59.921	58. 995	43. 190	30.786	51.04
HEMBA 1005513	167. 332	70.217	88.519	56.620	54. 920	73.797	80.751	68. 62 18. 61
HEMBA1005517	37.667	10, 443	23.901	8.903	17.777	21.966	23.844	27.82
HEMBA1005518	109. 105	25.679	71, 345	23.319	36.856	47.397		94.08
			100 400		1 400 007	AC 670	1 67 100	
HEMBA 1005520	200. 267	104.176	459.373	133. 255	106.207	95.070	67.199	
HEMBA 1005520 HEMBA 1005522	36. 421	15.946	24.796	12.598	8. 472	14.558	16.899	13.85
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526	36. 421 116. 274	15. 946 72. 899	24. 796 292. 397	12,598 82.002	8. 472 73. 603	14.558	16.899 34.319	13.85 47.68
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528	36. 421 116. 274 13. 037	15.946 72.899 9.406	24. 796 292. 397 30. 550	12.598 82.002 14.612	8. 472 73. 603 15. 947	14. 558 66. 198 16. 516	16.899 34.319 7.583	13.85 47.68 24.98
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005530	36. 421 116. 274 13. 037 56. 516	15. 946 72. 899 9. 406 26. 583	24. 796 292. 397 30. 550 63. 811	12,598 82,002 14,612 13,686	8. 472 73. 603 15. 947 21. 441	14, 558 66, 198 16, 516 29, 159	16.899 34.319 7.583 24.254	13.85 47.68 24.98 21.71
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005530 HEMBA 1005530	36. 421 116. 274 13. 037 56. 516 5. 523	15.946 72.899 9.406 26.583 17.373	24. 796 292. 397 30. 550 63. 811 36. 952	12.598 82.002 14.612 13.686 7.017	8. 472 73. 603 15. 947 21. 441 10. 885	14, 558 66, 198 16, 516 29, 159 11, 406	16.899 34.319 7.583 24.254 15.411	13. 85 47. 68 24. 98 21. 71 35. 78
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005530 HEMBA 1005530 HEMBA 1005539	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424	12.598 82.002 14.612 13.686 7.017 17.584	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989	14. 558 66. 198 16. 516 29. 159 11. 406 35. 829	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772	13. 85 47. 68 24. 98 21. 71 35. 78 25. 91
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005538 HEMBA 1005539 HEMBA 1005539	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124	12.598 82.002 14.612 13.686 7.017 17.584 15.206	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822	14.558 66.198 16.516 29.159 11.406 35.829 33.595	16.899 34.319 7.583 24.254 15.411 28.772 31.865	13. 85 47. 68 24. 98 21. 71 35. 78 25. 91 24. 09
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005538 HEMBA 1005538 HEMBA 1005539 HEMBA 1005545 HEMBA 1005545	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050	12.598 82.002 14.612 13.686 7.017 17.584 15.206 10.139	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638	14.558 66.198 16.516 29.159 11.406 35.829 33.595 22.115	16.899 34.319 7.583 24.254 15.411 28.772 31.865 22.271	13. 85 47. 68 24. 98 21. 71 35. 78 25. 91 24. 09 39. 29
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005530 HEMBA 1005530 HEMBA 1005533 HEMBA 1005534 HEMBA 1005548 HEMBA 1005548	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831	12,598 82,002 14,612 13,686 7,017 17,584 15,206 10,139 84,934	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638 81. 893	14.558 66.198 16.516 29.159 11.406 35.829 33.595 22.115 79.223	16.899 34.319 7.583 24.254 15.411 28.772 31.865 22.271 60.281	13.85 47.68 24.98 21.71 35.78 25.91 24.03 39.29 62.08
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005526 HEMBA 1005530 HEMBA 1005530 HEMBA 1005530 HEMBA 1005538 HEMBA 1005538 HEMBA 1005535 HEMBA 1005535 HEMBA 1005553	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021	24. 795 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831 24. 397	12, 598 82, 002 14, 612 13, 686 7, 017 17, 584 15, 206 10, 139 84, 934 9, 638	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638 81. 893 22. 919	14. 558 66. 198 16. 516 29. 159 11. 406 35. 829 33. 595 22. 115 79. 223 24. 422	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772 31. 865 22. 271 60. 281 21. 466	13.85 47.68 24.98 21.71 35.78 25.91 24.09 39.29 62.08
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005530 HEMBA 1005530 HEMBA 1005530 HEMBA 1005538 HEMBA 1005548 HEMBA 1005548 HEMBA 1005558 HEMBA 1005558 HEMBA 1005568	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488 74. 152	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021 61. 206	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 15. 050 363. 831 24. 397 184. 989	12, 598 82, 002 14, 612 13, 686 7, 017 17, 584 15, 206 10, 139 84, 934 9, 638 53, 681	8. 472 73. 603 15. 947 21. 441 10. 385 24. 989 46. 822 18. 638 81. 893 22. 919 38. 261	14, 558 66, 198 16, 516 29, 159 11, 406 35, 829 33, 595 22, 115 79, 223 24, 422 33, 077	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772 31. 865 22. 271 50. 281 21. 466 24. 038	13.85 47.68 24.98 21.71 35.78 25.91 24.09 39.29 62.08 8.17
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005526 HEMBA 1005530 HEMBA 1005530 HEMBA 1005530 HEMBA 1005535 HEMBA 1005545 HEMBA 1005546 HEMBA 1005566 HEMBA 1005568 HEMBA 1005568	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488 74. 152	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021 61. 206 68. 747	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831 24. 397 184. 989 74. 768	12. 598 82. 002 14. 612 13. 686 7. 017 17. 584 15. 206 10. 139 84. 934 9. 638 53. 681 17. 273	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638 81. 893 22. 919 38. 261 26. 562	14, 558 66, 198 16, 516 29, 159 11, 406 35, 829 33, 595 22, 115 79, 223 24, 422 33, 077 31, 212	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772 60. 281 21. 466 24. 038 27. 080	13. 85 47. 68 24. 98 21. 71 35. 78 25. 91 24. 09 39. 29 62. 08 8. 1 37. 01
HEMBA 1005520 HEMBA 1005526 HEMBA 1005526 HEMBA 1005526 HEMBA 1005530 HEMBA 1005530 HEMBA 1005533 HEMBA 1005534 HEMBA 1005548 HEMBA 1005548 HEMBA 1005568 HEMBA 1005560 HEMBA 1005560 HEMBA 1005570 HEMBA 1005570	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488 74. 152 54. 151 71. 454	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021 61. 206 68. 747 57. 260	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831 24. 397 184. 989 74. 768	12. 598 82. 002 14. 612 13. 686 7. 017 17. 584 15. 206 10. 139 84. 934 9. 638 53. 681 17. 273 21. 283	8, 472 73, 603 15, 947 21, 441 10, 885 24, 989 46, 822 18, 638 81, 893 22, 919 38, 261 26, 562 8, 931	14, 558 66, 198 16, 516 29, 159 11, 406 35, 829 33, 595 22, 115 79, 223 24, 422 33, 077 31, 212 30, 461	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772 31. 865 22. 271 60. 281 21. 466 24. 038 27. 080 29. 371	13.85 47.68 24.98 21.71 35.78 25.91 24.09 39.29 62.08 8.17 37.01 30.22
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005528 HEMBA 1005538 HEMBA 1005539 HEMBA 1005539 HEMBA 1005543 HEMBA 1005545 HEMBA 1005548 HEMBA 1005558 HEMBA 1005558 HEMBA 1005576 HEMBA 1005576	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488 74. 152 54. 151 71. 454	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021 61. 206 68. 747 57. 260	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831 24. 397 184. 989 74. 768 39. 016	12. 598 82. 002 14. 612 13. 686 7. 017 17. 584 15. 206 10. 139 84. 934 9. 638 53. 681 17. 273 21. 283	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638 81. 893 22. 919 38. 261 26. 562 8. 931 6. 610	14, 558 66, 198 16, 516 29, 159 11, 406 35, 829 33, 595 22, 115 79, 223 24, 422 30, 461 30, 461 18, 266	16. 899 34. 319 7. 583 24. 254 15. 411 28. 772 31. 865 22. 271 60. 281 21. 466 24. 038 27. 080 29. 371 12. 838	13. 85 47. 68 24. 98 21. 71 35. 78 25. 91 24. 09 39. 29 62. 08 8. 17 37. 01 30. 22 19. 99
HEMBA 1005520 HEMBA 1005522 HEMBA 1005526 HEMBA 1005526 HEMBA 1005530 HEMBA 1005530 HEMBA 1005533 HEMBA 1005545 HEMBA 1005545 HEMBA 1005546 HEMBA 1005566 HEMBA 1005566 HEMBA 1005566 HEMBA 1005570 HEMBA 1005570	36. 421 116. 274 13. 037 56. 516 5. 523 76. 498 46. 912 57. 779 141. 489 52. 488 74. 152 54. 151 71. 454	15. 946 72. 899 9. 406 26. 583 17. 373 30. 847 10. 940 14. 326 120. 695 20. 021 61. 206 68. 747 57. 260 13. 448 27. 270	24. 796 292. 397 30. 550 63. 811 36. 952 69. 424 32. 124 15. 050 363. 831 24. 397 184. 989 74. 768 39. 016 21. 181 38. 708	12.598 82.002 14.612 13.686 7.017 17.584 15.206 10.139 84.934 9.638 53.681 17.273 21.283 13.021	8. 472 73. 603 15. 947 21. 441 10. 885 24. 989 46. 822 18. 638 81. 893 22. 919 38. 261 26. 562 8. 931 6. 610	14, 558 66, 198 16, 516 29, 159 11, 406 35, 829 33, 595 22, 115 79, 223 24, 422 30, 461 30, 461 18, 266	16. 899 34. 319 7. 583 24. 254 25. 411 28. 772 31. 865 22. 271 60. 281 21. 466 24. 038 27. 080 29. 371 12. 838 28. 804	13. 85: 47. 68: 24. 98: 21. 71: 35. 78: 25. 91: 24. 09: 39. 29: 62. 08: 8. 17: 37. 01: 30. 22: 19. 99: 10. 18: 16. 84:

Table 34

	DENATA AREFAA	00 764 1	00 100 1	00 630	0 725 1	10.545	12.468	10.523	17.884
	HEMBA1005583	23, 564	22. 466	98.629	9. 735				52.441
	HEMBA 1005588	96.041	96. 264	266.022	69. 126	54. 588	44. 105	34.310	
5	HEMBA 1005593	61.102	40. 350	125.688	37. 987	35. 953	41.577	39.834	47.357
5	HEMBA1005595	52, 429	18.652	31. 240	8.095	8.750	14.586	12.433	7.264
	HEHBA1005597	125.119	43. 335	90.414	24. 402	44.780	74.946	66.352	45. 322
	HEMBA1005606	141.646	66.667	95. 041	30.084	57. 974	107.329	84.655	46, 145
					52.002	41.602	40.406	26. 928	42, 614
	HEMBA1005609	77. 991	60. 190	244. 951					
	HEMBA1005616	47.760	62.865	190.870	42.670	41.809	32. 256	23.683	43. 139
10	HEMBA 1005621	33.797	18. 993	22. 515	11.333	11. 545	16.964	12.122	13. 910
10	HEMBA1005627	128.661	66. 487	148. 021	45. 359	42.161	42.054	30.884	43.319
	HEMBA1005628	43.539	36, 758	85.714	25. 524	46.601	19. 229	82.784	36. 636
	HEMBA 1005631	21, 340	8. 467	38.068	22.476	18. 318	17.813	12.599	28. 199
	HEMBA1005632	113.190	73.661	233.637	59.097	45. 388	52.090	29.944	37. 461
		123.668	195. 912	390. 579	101. 523	107. 528	72.729	54.939	130. 473
	HEMBA 1005634				7. 453	5. 561	13.084	8.973	5. 282
15	HEMBA1005662	15.391	11.345	23.021				13.796	28. 043
,,	HEMBA1005666	33.844	30, 419	34. 983	13.220	31. 573	24. 609		
	HEMBA 1005670	91.667	63.609	255. 523	57.730	46. 927	45. 285	23.794	46.684
	HEMBA 1005671	63.448	55. 388	34. 948	26. 297	20. 567	2.367	5.666	13.509
	HEMBA1005679	53.089	33. 284	126. 705	39.666	32. 151	40.446	37.522	36.817
	HEMBA1005680	115. 289	72.018	220. 408	76.653	55. 707	68.735	32.613	36. 282
	HEMBA1005685	68.783	45.211	72.197	30.110	32.724	43.022	37.740	33. 510
20	HEM8A1005698	37.890	35.679	44.793	29.794	25. 150	48.613	15.651	20.648
	HEMBA1005699	14. 243	17. 539	37.269	9.035	12. 276	5. 454	5, 259	6.787
	HEP8A1005703	19.524	15.116	20.249	7,662	15. 489	11.648	8.488	10, 229
	HEM8A1005705	35.316	35.677	66.552	26. 492	29.605	90. 298	18.303	44.730
				30. 267	17. 829	17.668	17.695	14.517	23.820
	HEMBA1005712	20.312	29.695			13.391	32.044	15.084	7.078
	HEMBA1005717	47.313	15.037	30.499	6. 950				76.705
25	HEMBA1005718	88, 576	81.734	176.773	75.414	46.080	58.797	49.803	
	HEMBA 1005721	84. 981	42.340	58. 434	18. 134	34. 246	43. 284	34.523	41.450
	HEMBA1005722	174.952	92. 346	194.868	55.652	48.768	63. 471	92.755	56.031
	HELMA1005724	32.655	8. 284	5. 342	4.000	14.801	15. 671	9.324	5. 953
	HEMBA1005732	89.624	24.907	32.546	5.638	21.753	30.046	28. 487	20.595
	HEMBA1005737	25. 179	16.797	16.017	10.703	12.731	12.444	8.579	7.257
	HENBA1005742	11.547	23.162	24. 345	20.921	29. 934	18.597	13.749	22.702
30	HEBBA1005748	36.098	14.407	21.907	16.923	13. 431	12.235	10.908	8.606
	HEMBA1005747	80.718	30.396	44. 843	21.861	30. 274	80, 588	47.082	26.037
	HEMBA1005749	35.749	31.758	64.769	22.766	28.853	26.733	31.698	30.753
				30.663	37.837	21.308	24. 392	15. 905	25.470
	HEMBA 1005755	34.680	39. 133			28. 933	46. 295	36. 173	31.205
	HEMBA1005760	118. 125	41.490	33.276	25. 724			31.098	40.041
35	HEMBA 1005765	94. 451	70.516	200.826	48.023	37. 340	35.414		
33	HEMBA 1005765	112.861	70.359	87.247	48. 958	51.073	52.147	72.391	63. 859
	HEMBA1005780	55, 961	34.713	89.816	28. 466	46.254	28. 283	25.156	29. 122
	HEMBA 1005795	18.800	38. 386	19.666	10, 007	13.009	11.811	13.106	14. 493
	HEMBA 1005809	67.301	66.510	87.390	53.061	43. 975	35. 574	35. 334	57.818
	HEMBA1005813	52.911	84.881	160.064	38.752	43.727	30.799	23. 426	57.177
	HEMBA1005815	30, 398	30, 434	43.366	19.911	16.123	39.746	26.743	28. 548
40	HEMBA1005822	40.948	47.746	65.298	51.932	30.845	20. 187	22.641	29.114
	HEMBA1005829	114.982	70.536	272.004	48.816	36.558	40.259	23.443	35. 824
	HEMBA1005833	59, 540	25.743	29. 266	15. 545	24.711	26.964	17.968	18.807
	HEMBA1005834	151, 440	82.917	322.413	102.348	74.711	59, 590	35.082	70.415
	HEMBA1005844	66.624	11.865	96. 556	95.719	56.133	75. 546	55. 974	122.840
						85.623		90.291	87.310
	HEMBA 1005852	71.743	77.830	1 ,				27. 359	58. 468
45	HEMBA1005853	62.809	83. 326	343. 381	63.897	79. 208	48. 939		
	HEMBA1005878	139, 991	109. 928	447.600	93.748	65. 325	53.917	35. 383	63.446
	HEMBA1005883	5, 211	6.310	6.808	14. 769	10.070		4.485	11.850
	HEMBA 1005804	9.136	10.768	29. 442	9. 504	7. 302	9. 142	4.561	12.287
	HEMBA1005891	8. 927	12,500	12,662	5. 996	7.370	7.346	1.250	5. 470
	REMBA 1005894	70,006	59.347	177.879	49. 407	29. 584	23. 227	14.651	36.934
	HEMBA1005898	84, 399	61. 254	234. 549	59.872	43. 955		23.019	41.130
50	HEI*BA1005902	38.306	16.873	52.804	16.742	33.189		26.778	43.681
				8.804	5. 339	3. 957	17.078	5.311	4. 941
	HEPBA1005907	4.806	3. 997	23. 479			6. 683	0,750	10.643
,	HEMBA1005909	4.140	3.733	316 300	42 107		<del></del>	39.302	55. 189
	HEL 8A1005911	143. 926	92.633	316.302					26.318
	HEMBA1005912	18.801	17. 269	13.568	32. 298	21.976	14.454	12.917	20.318

Table 35

	HEHBA1005913	10. 533	15.117	14.368	16.655	8. 179	7.135	7.907	12.918
	HEHBA1005921	83. 262	45. 648	252.573	51.044	41.764	22. 286	23.762	46, 202
5	HEBBA1005922	64. 440	17.427	35. 136	20.084	33.779	24.835	18. 394	14.883
3	HEMBA1005929	173.002	139.696	378.444	96.543	83.075	72.298	55. 205	94.716
	HEMBA1005931	146.354	89.551	224.601	83.623	63.406	73.122	54. 973	59.891
	HEMBA1005934	141.558	91.791	227.012	89.834	99.341	96.876	62.967	55. 492
	HEMBA1005945	144.693	21.871	38, 980	19.915	46.699	78. 590	80.430	30.052
	HEMBA1005962	67, 209	34.719	63,745	21.004	17, 931	29. 331	21.199	20.008
	HEMBA1005963	18. 320	6.954	9, 127	5.913	2.497	8.674	7.674	4. 873
10	HEMBA1005990	581.646	117. 336	139.967	53.671	242.262	424. 182	418.873	85. 511
	HEMBA1005991	67.437	59.327	188.570	42.994	21, 101	33.868	19. 164	11.619
	HEMBA1005999	193, 878	135. 695	450.789	126. 399	129, 150	103.289	53. 193	115. 911
	HEMBA1006002	73.560	25. 438	22.156	12.657	16,731	16.116	10, 600	19. 305
	HEMBA1006005	59.620	7.083	16.863	8.213	29.019	53.513	52.130	23.838
	HEMBA1006011	25. 811	30, 413	39.888	21.434	54, 488	30.978	27. 996	25. 339
15	HEMBA1006013	51.604	13. 251	19.743	11.817	15.364	28.363	21.493	18.674
	HEMBA1006016	101.929	42.149	115.996	36.228	39.875	45.607	33, 305	26. 397
	HEMBA1006019	31.772	18. 482	22.979	15. 207	22.984	24.244	26. 246	14. 100
	HEMBA1006021	26. 984	10.213,	45. 937	9.253	20.615	14. 587	14. 203	12.296
	HEMBA1006022	100. 930	40.046	62.368	42.744	23.660	46.057	25.008	19. 323
	HEMBA1006031	42.088	41.281	14.729	11.264	12.725	36.716	13.037	5. 133
20	HEMBA1006035	10.089	10.059	27.290	8. 123	6.309	6.629	2. 039	5. 229
	HEMBA1006035	188. 431	82.469	443.914	119.939	80.135	81.126	54.157	94.631
	HEMBA1006042	69. 906	33.773	134. 462	30.108	23.244	29.765	29.479	29. 607
	HEMBA1006044	53.721	10.199	12.818	4.725	8.467	5. 436	2.586	4. 088
	HEMBA1006045	48.078	43.730	61.128	28. 336	25. 311	26.451	23. 478	44. 272
	HEMBA1006048	35. 685	18.435	41.495	19. 225	19.636	34. 213	26.302	28. 809
25	HEMBA1006053	0.000	356.500	78.844	24. 270	47.030	114.986	63. 574	385. 970
23	HEMBA1006055	7.603	5. 331	12.625	4. 484	13.776	12.227	9.079	5. 545
	HEMBA1006058	51.872	19.394	14.828	7.834	11.877	25. 640	15, 830	21.486
	HEMBA1006053	72.886	52.429	63.882	34.021	30.125	39. 536	28. 303	35.860
	HEMBA1006067	6.005	14.253	7.505	3. 169	2.242	3.352	4. 358	0.888
	HEE8A1006081	70. 282	19.151	25.838	8.981	9.908	26. 560	16.837	23.976
20	HEMBA1006089	54. 392	23.145	42.709	18. 278	17.433	17.768	18. 372	23. 981
30	HEMBA1006090	71.092	20. 389	36.832	15. 386	17.868	38.904	35. 031	18. 238
	HEMBA1006091	59. 022	28. 947	126. 425	16.353	30.302	55.034	53.660	66.468
	HEMBA1006093	111.885	11.435	50.738	16. 185	27.687	43.178	26.048	14. 980
	HEMBA1006099	40.381	27. 136	39.149	18. 199	31.100	31.158	28. 536	25. 484
	HEMBA 1006 100	36, 979	48. 991	259. 267	41.090	50.094	24.833	13.379	34. 456
0.5	HEMBA 1006 108	40. 170	19. 301	21.811	11. 125	8.795	12.441	8.780	16. 453
35	HEMBA1006114	42.849	44.783	46.702	33. 193	23. 220	34. 626	28. 294	51.756
	HEMBA1006121	160. 208	21.943	26.728	10.160	21.331	17. 129	26.838	25. 137
	HEMBA1006124	63.151	11.764	15.994	17.764	14.099	57. 249	29. 200	8. 240 68. 939
	HEMBA 1006 125	72.730	70. 406	57.020	50.057	45. 287	40.856	45. 665 25. 726	21.315
	HEKBA1006130	36.221	31.688	34.742	7.817	28. 246	34.473	67, 854	89.604
	HERBA1006138	160. 258	170.815	435. 120 238. 562	106.719 54.531	139,660	65.032	45. 938	59.791
40	HEMBA 1006 142 HEMBA 1006 150	127. 194	85. 725 58. 231	76.666	59. 941	19.605	46.114	33. 261	75, 731
	HEMBA 1006 151	66.777 189.265	57. 959	104.921	29.646	46. 546	66.736	74, 155	88. 383
	HEMBA1006155	141. 288	19. 560	50, 142	11.752	32.711	79. 435	60.621	32.838
	HEMBA1006158	17.276	12.039	19.210	7. 139	7.468	23.241	7.360	13.357
	T		<del></del>		97. 488	87.832	69.460	42.210	85. 135
	HEMBA1006164	66.839	70.843	382.965	13. 911	21.700	40.783	26.049	37. 233
45	HEMBA1006173	63.939	35. 393	52. 598	22. 894	32.403	35.413	40.872	67.870
	HEL BA1006176	51.671	222.661	52.703	39. 369	29. 305	59.271	24.272	83.343
	HEPBA1006182	72.842	38. 362	132. 455	29. 730	26.735	30.382	19, 907	34. 405
	HEMBA1006197	16.655	31. 338	37. 528	55. 808	23. 143	18.848	13.456	40.765
•	HELBA1008198	30.466	15. 178	21. 337	16. 185	25.764	15.643	14. 389	18.561
	HELBA1006213	38.783	20. 120	38. 136	15. 627	10.604	25. 761	21.716	35. 282
50	HEMBA1006217	32,003	18. 510	33.960	4. 079	17. 107	31.016	36.526	19.419
	HEMBA1006226	40.304	60.090	110. 529	40.359	39.915	62.796	35. 202	59. 281
	HEMBA1006235	40.954	9.021	21.361	7.280	14. 241	13.056	4. 951	7.077
	HEMBA1006248	42.946	17. 521	32.092	10.747	12.992	19. 331	18. 339	17. 999
	HEMBA1008251	84.944	24. 303	30. 554		24.212	30.870	18.154	10.996
		· · · · · · · · · · · ·							

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Table 36

							-		77 777
	HEMBA1006252	36.069	24.512	74. 170	29.506	28.055	19.517	14.085	15.356
	HEMBA1006253	75.854	7.002	20.773	16. 455	11.705	12.936	6,506	11.398
•	HEMBA1006259	37.456	48. 402	136,000	39.735	19.462	25. 242	19.832	25. 931
5	HEMBA1006261	23.677	23. 578	5.874	13.012	7.127	59. 427	7.141	17. 143
	HEMBA 1006268	35.886	12.563	30.879	8.970	7.077	19.793	22. 288	18. 289
	HEMBA1005271	122, 980	98.618	185.469	77.610	45.268	47.910	36. 533	47. 867
	HEMBA1006272	16, 261	12.829	9, 416	4.968	5.925	27.766	15.997	7. 567
	HEMBA1006273	47.890	12.641	71.219	20.880	30.446	30.473	18. 419	22.459
	HEMBA1006276	79. 296	11.878	30.854	34. 032	8.760	27.168	16.165	7. 501
10	HEMBA1006278	40. 093	7.717	26.091	4.506	18.669	11.680	11. 224	9.893
	HEMBA1006283	16. 994	23. 586	25.614	25. 226	23.447	59.086	28. 267	25.848
	HEMBA1006284	29. 982	22. 166	27.891	20.874	8, 594	18. 386	15. 293	13. 396
	HEMBA1006291	22.745	13.071	36.861	9.670	4.059	11.649	31.851	7.519
		17.718		20. 081	10.169	4, 378	7. 903	9. 259	7.898
,	HEMBA1006292 HEMBA1006293	31.307	8. 916 10. 056	8.749	4.645	4. 097	6.631	8. 473	7. 189
15					1.371	4. 543	2. 465	1.701	2. 648
7.5	HEMBA 1006299	21.091	5. 917	6, 157			31.160	24. 850	17.764
	HEMBA1006309	69.975	25. 568	110.869	33. 191	19.510		27. 263	17.748
	HEMBA1006310	40.983	23, 265	36. 585	20.570	11.748	29.056		19. 313
	HEMBA1006311	85. 398	20.844	64.711	8. 925	20. 171	92.798	9. 481	5. 082
	HEMBA1006313	27.762	12.975	47.707	17.417	7. 455			3. 270
	HEMBA 1006316	23. 345	3, 751	3. 303	2.158	8.774	9.668	8. 505	85.968
20	HEMBA1006328	79. 937	83.744	185. 981	41.111	28.820	37.527	35. 377	
	HEMBA1006334	22.524	16.717	17.679	5.994	8. 506	9.813	3. 866	5.361
	HEMBA 1006335	72.666	41.477	35. 235	27. 435	6.110	5.851	24. 375	8. 434 25. 330
	HEMBA 1006344	34.707	67.866	132. 978	46.518	34.812	40.158	41. 934	
	HEMBA1006347	34. 301	16.445	32.190	19.603	15.749	20.762	20. 884	15.376
	HEMBA1006349	139.389	26.300	48. 767	43. 275	22.026	24.648	22.876	21. 499
25	HEMBA1006352	21.127	17.873	15. 526	9.410	8.472	14.845	7. 491	9.414
	HEMBA1006357	94, 337	82.319	287. 531	67.888	76. 120	47.179	41.500	59.557
	HEMBA 1006358	48. 925	31. 345	132.494	32.473	25.019	28. 197	13. 250	24. 899
	HEMBA 1008359	57. 203	18. 522	160. 314	70.923	17. 441	30.686	11.154	47. 991
	HEMBA 1006360	29.518	10. 133	15, 515	17. 275	6. 141	13.876	6.804	8. 361
	HELBA1006354	59. 236	7. 900	27. 522	12, 114	5. 401	15. 432	17. 981	6. 672
30	HEMBA1005377	67.120	31.113	57. 269	33.567	23.849	45. 246	31.609	20. 280
50	HEMBA 1006380	73.227	57. 029	182. 581	57.870	22. 288	33.416	23.616	40. 932
	HEMBA 1006381	359. 346	122. 755	376.090	126.304	112.826	146. 346	91. 469	93. 252
	HEMBA 1006385	60.234	62. 166	257. 945	59.429	59. 157	40.136	35. 385	17. 281
	HEMBA 1005390	71.393	38.752	46.828	25.848	16. 455	41.253	16.013	27.609
	HEMBA1006391	61.261	18.765	20. 686	10.972	10.022	39. 431	27. 305	11.797
	HEMBA1006398	42.089	3. 225	18. 036	5. 299	25. 386	6.480	0.000	3.308
35	HEMBA1006405	137.413	28.645	40. 904	17.896	18. 180	84. 926	41. 325	24.773
	HEMBA1006410	149.580	32.840	61.022	20.027	39.718	54. 551	23.826	33.928
	HEMBA1006416	96.031	52.892	198.896	50. 538	38. 551	37.025	37. 809	33.447
	H£MBA1006418	23. 236	18. 335	23. 851	11.378	10. 280	28. 208	46. 245	36. 223
	HEMBA1006419	189. 293	101.979	476. 145	90.626	79.213	64. 306	40.042	52.384
	HEMBA1006421	39.702	26. 487	127. 221	23.773	16.184	14.460	12. 270	13.523
40	HE199A1008424	4. 484	35. 452	10. 588	3.778	4. 512	7.346	2. 324	3. 323
	HEMBA 1006426	88.597	67. 224	230.530	60.836	32. 273	40.489	17, 284	36. 244
	HEMBA1008430	61.672	17. 989	69. 151	15, 913	11.038	15. 595	9.696	17. 632
									11.056
	HEMBA1006438	45. 084	34. 475	111.512	27.012	15.035	34.111	12.678	
	HEMBA 1005438 HEMBA 1005445		13.919	53. 981	27.012 9.326	15. 035 15. 672	34.111	27.442	18. 331
	HEMBA1006438	45. 084			9. 326 1. 568	15.672 4.341	34. 167 2. 585	27. 442 1. 331	18.331 0.000
45	HEMBA1006448 HEMBA1006446 HEMBA1006456	45. 084 48. 245 22. 911 36. 915	13. 919 3. 160 28. 165	53. 981 3. 324 141. 114	9. 326 1. 568 18. 927	15. 672 4. 341 65. 823	34. 167 2. 585 33. 549	27. 442 1. 331 13. 651	18. 331 0. 000 33. 405
45	HEMBA1006445 HEMBA1006446 HEMBA1006456 HEMBA1006456	45. 084 48. 245 22. 911 36. 915 60. 747	13.919 3.160 28.165 42.392	53. 981 3. 324 141. 114 161. 108	9. 326 1. 568 18. 927 40. 447	15.672 4.341 65.823 22.274	34.167 2.585 33.549 32.823	27. 442 1. 331 13. 651 18. 018	18.331 0.000 33.405 27.165
45	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008466 HEMBA1008461 HEMBA1008467	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357	13.919 3.160 28.165 42.392 6.130	53. 981 3. 324 141. 114 161. 108 15. 734	9. 326 1. 568 18. 927 40. 447 10. 759	15.672 4.341 65.823 22.274 4.032	34. 167 2. 585 33. 549 32. 823 4. 471	27. 442 1. 331 13. 651 18. 018 6. 183	18. 331 0. 000 33. 405 27. 165 2. 655
45	HEMA1005438 HEMA1006445 HEMA1006446 HEMA1006456 HEMA1006461 HEMA1006467 HEMA1006467	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235	15.672 4.341 65.823 22.274 4.032 29.870	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286
45	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008466 HEMBA1008461 HEMBA1008467	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357	13.919 3.160 28.165 42.392 6.130	53. 981 3. 324 141. 114 161. 108 15. 734	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933	15. 672 4. 341 65. 823 22. 274 4. 032 29. 870 2. 522	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224	27.442 1.331 13.651 18.018 6.183 33.818 10.020	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873
45	HEMA1005438 HEMA1006445 HEMA1006446 HEMA1006456 HEMA1006461 HEMA1006467 HEMA1006467	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235	15.672 4.341 65.823 22.274 4.032 29.870 2.522 9.614	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875	27.442 1.331 13.651 18.018 6.183 33.818 10.020 17.655	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491
	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008456 HEMBA1008467 HEMBA1008467 HEMBA1008470 HEMBA1008474 HEMBA1008474	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498 43. 462	15. 672 4. 341 65. 823 22. 274 4. 032 29. 870 2. 522 9. 614 42. 248	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945
<b>45</b> <b>50</b>	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008456 HEMBA1008461 HEMBA1008467 HEMBA1008470 HEMBA1008470 HEMBA1008474 HEMBA1008476 HEMBA1008476	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504 12. 420 91. 936 169. 312	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381 63. 588	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498	15.672 4.341 65.823 22.274 4.032 29.870 2.522 9.614 42.248 57.839	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725 95. 521	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725 75. 480	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945 239. 325
	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008456 HEMBA1008461 HEMBA1008467 HEMBA1008470 HEMBA1008470 HEMBA1008474 HEMBA1008476 HEMBA1008476	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718 180. 042 129. 627	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504 12. 420 91. 936 169. 312	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498 43. 462	15. 672 4. 341 65. 823 22. 274 4. 032 29. 870 2. 522 9. 614 42. 248	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725 75. 480 23. 818	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945 239. 325 27. 130
	HEMBA1008438 HEMBA1008445 HEMBA1008446 HEMBA1008456 HEMBA1008467 HEMBA1008467 HEMBA1008470 HEMBA1008474 HEMBA1008474	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718 180. 042 129. 627 99. 620	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504 12. 420 91. 936	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381 63. 588 167. 982	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498 43. 462 151. 338	15.672 4.341 65.823 22.274 4.032 29.870 2.522 9.614 42.248 57.839	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725 95. 521	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725 75. 480	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945 239. 325
	HEMBA1008438 HEMBA1008445 HEMBA1008456 HEMBA1008456 HEMBA1008467 HEMBA1008470 HEMBA1008470 HEMBA1008474 HEMBA1008474 HEMBA1008476 HEMBA1008476 HEMBA1008482 HEMBA1008483	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718 180. 042 129. 627 99. 620 41. 690	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504 12. 420 91. 936 169. 312 64. 773 4. 055	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381 63. 588 167. 982 232. 207	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498 43. 462 151. 338 50. 445	15. 672 4. 341 65. 823 22. 274 4. 032 29. 870 2. 522 9. 614 42. 248 57. 839 29. 074	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725 95. 521 37. 572 9. 351 21. 229	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725 75. 480 23. 818	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945 239. 325 27. 130 10. 066 15. 706
	HEPBA1005438 HEPBA1006445 HEPBA1006445 HEPBA1006456 HEPBA1006457 HEPBA1006470 HEPBA1006471 HEPBA1006471 HEPBA1006474 HEPBA1006474 HEPBA1006483 HEPBA1006483	45. 084 48. 245 22. 911 36. 915 60. 747 13. 357 73. 960 19. 032 25. 718 180. 042 129. 627 99. 620	13. 919 3. 160 28. 165 42. 392 6. 130 30. 706 4. 504 12. 420 91. 936 169. 312 64. 773	53. 981 3. 324 141. 114 161. 108 15. 734 103. 625 7. 503 21. 381 63. 588 167. 982 232. 207 17. 445	9. 326 1. 568 18. 927 40. 447 10. 759 27. 235 2. 933 11. 498 43. 462 151. 338 50. 445 11. 682	15. 672 4. 341 65. 823 22. 274 4. 032 29. 870 2. 522 9. 614 42. 248 57. 839 29. 074 4. 522	34. 167 2. 585 33. 549 32. 823 4. 471 33. 756 5. 224 19. 875 109. 725 95. 521 37. 572 9. 351	27. 442 1. 331 13. 651 18. 018 6. 183 33. 818 10. 020 17. 655 88. 725 75. 480 23. 818 6. 411	18. 331 0. 000 33. 405 27. 165 2. 655 24. 286 1. 873 13. 491 65. 945 239. 325 27. 130 10. 066

Table 37

	CIEUS LANGE COA				26 451	0 026 [	0.026	11 410 1	12.090
	HEMBA1006492	14.002	19.916	24.662	35. 451	8.836 8.718	8.075	5.851	5. 887
	HEMBA1006494	7.279	0.000	19.790	3.750		8. 343		
5	HEMBA1006497	41. 284	12. 396	23. 326	6.590	7.186	11. 228	9.062	5. 781
Ū	HEMBA1006501	160.565	16.895	26.893	13.446	17.608	65. 467	41.560	6. 197
	HEMBA 1006502	53.451	19.114	39. 593	25.366	10.919	15. 054	17.536	15.658
	HEMBA1006507	19.274	8.180	10. 287	4. 521	7.939	5. 288	15. 480	10.062
	HEM8A1006517	95.989	30.085	91.871	18.732	21.918	45. 881	29.819	16.672
	HEMBA1006521	31.224	27.873	37. 864	18.318	9.774	14. 205·	14.646	13. 907
	HEMBA1006529	28.702	20.010	34.050	20.150	16. 588	7. 353	8.993	17. 327
10	HEMBA1006530	18.445	16.411	29. 175	14. 433	12. 214	16.734	15, 731	8.081
	HEMBA 1006535	11.627	7.208	18.048	3.956	8. 160	19.824	5. 837	3. 457
	HEMBA1006536	68.087	40.009	142.475	43, 263	34. 343	42.050	42. 157	23.975
	HEMBA1006540	20.393	10.867	35, 153	8.637	8. 656	15.027	11.094	10.350
	HEMBA1006544	30, 281	4. 662	59. 940	7.791	7.169	15.883	8.745	8.693
	HEMBA1006546	68.722	53, 155	127, 193	49. 337	73.807	60.506	22.328	34.045
15	HEMBA1006549	13. 885	13.666	21.800	11.666	8. 491	14, 211	8.987	6.080
	HEMBA1006559	26.976	22.040	38, 197	16.910	14.550	14.058	13.018	17.217
	HEMBA1006562	55. 924	24.663	75. 789	20.363	17. 181	26.651	18.158	19.510
	HEMBA1006566	20.849	6. 116,	14. 933	8.767	9. 572	6.937	5. 229	4.788
	HEMBA1006569	67.508	20. 299	44. 291	27.048	12.798	15. 243	24.739	31.861
	HEBBA 1006572	21.817	4. 339	15.862	1.796	3, 407	11.582	8. 381	5. 922
20	HEBBA1006579	5, 427	18. 336	4.219	3.440	2.139	5. 460	3.967	5.110
-0	HEMBA1006583	31.967	15. 854	29.307	14.271	11.747	26.889	17.058	10.451
	HEM8A1006595	59.014	41.577	148. 359	30.660	16.681	19. 571	13. 265	24.768
	HEMBA1006597	111.817	64. 480	210.001	47.574	27.392	47.009	27.887	28.666
	HEMBA1006606	79.184	47.311	131.822	40.177	33. 228	35. 403	25. 240	31.687
	HEMBA1008612	43. 105	20. 909	46.913	39.205	20. 348	25, 383	18.706	17. 150
	HEEBA1006617	79.139	62.924	235.236	60.258	30. 407	40.264	28. 184	38.643
25	HEMBA1006624	449. 384	84.050	165.494	39. 352	209, 908	291, 427	208.533	65. 478
	HEMBA1006631	168.309	108.316	381.778	89.696	71.812	80.634	39. 325	50.996
	HEMBA1006635	51.406	33.730	158. 286	28.605	19.347	19, 781	9.639	12.894
	HEMBA1006839	67.363	30. 354	51.867	15. 409	33.210	43.083	25. 295	12.985
	HEMBA1006843	229.685	30. 246	56.218	16.406	35.196	68. 642	41.724	17, 931
	HEMBA1006648	80.985	32.464	39.607	14. 926	36.718	12. 135	32.217	48.853
30	HEMBA1006652	118. 455	69. 232	231.917	50.609	51.023	50.716	21.698	29.527
	HEMBA1006653	46.971	16.614	46, 472	16.579	12.358	15.364	13.867	9. 224
	HEMBA1006658	89.823	28. 363	60.976	37.660	28. 124	47.014	33, 470	15.872
	HEMBA1006659	79.863	33.626	48.217	49. 132	29. 124	33.070	25. 182	33.784
	HEMBA1006665	25.726	26.740	39.661	13.975	13. 287	15. 240	12.046	10.419
•	HEMBA1006666	8. 276	4. 281	10, 565	6.319	4. 257	10.392	2.791	2.171
35	HEMBA1005671	39.553	178.623	135.413	18. 941	17. 294	37.782	10.166	32.048
••	HEMBA1006674	100.472	44.108	176.724	46.922	36.367	44, 809	43.576	43. 269
	HEMBA1006676	120.417	42.888	163.816	29. 504	40. 435	60.162	32.540	34. 825
	HEPBA1006682	27.104	2. 556	23. 174	4. 035	8. 982	19.092	3.958	0.000
	HEYBA1006688	57. 351	56. 288	111.358	60. 597	65. 322	37.545	20.757	20. 789
	HEPBA1006695	132.496	140. 334	315.655	97. 296	56. 206	54. 392	37.622	57.596
40	HEEBA1006596	65.136	25. 204	42. 137	26.654	26.490	30. 156	6.159	27.512
40	HELBA1006702	4. 275	4. 328	8.881	7.114	3.362	1.846	7.796	1.965
	HEPBA1008707	52.417	20.765	26.862	21.409	19.843	32.229	13.146	18.546
	HELBA1006708	126.875	38. 520	66.803	31.253	33.294	55. 347	32.071	18. 229
	HEI BA1006709	67.500	31.686	94, 432	24. 924	17, 365	30.329	18,603	23.474
	HEMBA1008717	110.641	21.536	29. 255	12.664	16.091	54.326	26.752	11.544
	HEMBA1006724	34, 421	23.073	25, 607	18.231	12, 305	27.570	10.585	17.797
45	HEMBA1005731	35. 072	18. 255	41, 441	15. 382	16.479	17.272	10.826	15.482
	HEMBA1006737	60.467	14. 107	30.096		20. 232	22.606	10.316	11.440
	HEMBA1006742	60. 258	45. 190	134.964		21.315	21.889	15.223	23.529
	HEMBA1006743	41.970	22.864	31.760		15, 126	23.989	13.179	16.281
	HEMBA1008744	181.068	97.273	433.004		69.785	59, 354	46.770	-61.806
			9.753	37, 994		23.164	34.516	28, 426	23. 238
50	HEYBA1006749	51.776	60.318	88.111		47, 490	69.451	37.541	47.074
	HEMBA1006752	124. 800	30.459			17.745	16, 269	10.783	12.424
	HEMBA1008754	49. 957	21.737	26, 190		18. 249	38, 492	30.654	15. 933
	HEMBA1006758	75. 460					13. 334	8. 382	8. 573
	HEMBA1006767	14. 002	15. 106	11.961	16.059			, 0.002	
	HEMBA1006770	120, 485	21.505	62.144	29. 559	32.512	49.739	45. 952	28.318

Table 38

	HEMBA1006779	81, 492	51.077	162.657	41.163	39, 166	36.722	18.025	29.256
	HEMBA1006780	78.359	78.052	345. 442	73.371	68.858	55. 888	41.524	39.494 -
									25, 701
5	HEMBA1006789	29. 455	21.233	20.440	14. 349	11.547	38, 549	19.736	
	HEMBA1006795	143.727	88, 701	218.732	55.068	49.500	46. 284	21, 141	40.750
	HEMBATOD6796	87.214	15, 814	115. 542	17.685	16.790	38, 694	15. 525	15.352
									30.733
	HEMBA1006805	68.116	31. 212	153.041	33. 162	30.301	34. 197	24. 275	
	HEMBA1006807	94. 524	86.723	157. 559	64. 349	36. 505	62.933	23.097	55. 508
•	HEMBA1006813	40.696	4, 415	4.750	4. 264	10.978	7,562	6. 201	3.198
						17.006	30.866	20. 346	6, 250
10	HEMBA1006819	53.717	15. 217	30.071	14.679				
• •	HEMBA1006821	39.052	30.425	111.325	35.769	34. 975	22.216	18. 924	20.698
	HEMBA1006824	68, 491	61.498	201.721	47, 107	40. 322	27. 255	21.689	27.074
				102.038	77.046	40. 147	75. 996	66.799	71,706
	HEMBA1006832	84. 462	89.500						
	HEMBA1006834	123.958	57. 085	160.407	48. 909	41.460	61.443	30.402	31, 940
	HEMBA1005835	33,705	19, 529	38. 470	23. 193	18.979	22.344	22. 426	16.742
	HEM8A1006843	52, 436	44. 642	96.773	258.615	195.878	33.141	8. 256	13.117
15							34. 943	15.743	28. 240
	HEMBA1006849	88. 931	34. 224	158.388	39.483	30.349			
	HEMBA1006850	44.733	24. 923	67.667	24. 186	15.829	36.593	11.223	18. 454
	HEMBA1006861	215. 207	94. 180	158.997	67.349	259.512	135.856	371.932	44.063
			59.773,	124. 376	43.328	69.356	71.072	66.350	45. 129
	HEMBA1006865	124. 996							
	HEMBA1006867	16.632	11,094	39.646	14.084	12. 902	11.855	5.865	18.338
	HEMBA1006873	9.965	9, 279	7.010	5.013	6.262	5. 127	7.141	8.422
20	HEMBA1006877	44.043	18. 321	20.546	8. 172	14.670	13. 165	16.493	9, 073
-					25.739	29. 525	48.800	41.513	17. 905
	HEMBA1006878	100. 427	34. 418	109.029					
	HEMBA1005879	108. 299	42.811	121.051	60.872	47.507	40.075	14. 429	51. 924
	HEMBA1006884	95. 426	29, 331	67.556	27.787	25.909	106.818	47.878	47.793
	HEMBA1006885	107, 720	54. 342	127, 920	62.272	55.739	51.739	36.790	50, 612
								26.952	19.149
	HEMBA1006886	50. 841	22.970	51. 528	12.561	20.660	23. 207		
25	HEMBA1006889	81.809	20. 952	21.474	12.691	24.681	41.822	48.768	15. 196
	HEMBA1006896	68. 030	97. 285	75.370	52.746	23.109	44. 481	37.701	50.662
	HEMBA1006900			61.016	23.329	21.390	38.404	27.583	22.774
		61.515	36.410						
	HEMBA1006902	43. 283	19.713	47. 129	12.105	11.602	27.830	26.548	13. 885
	HEMBA1006912	183.904	90, 995	338. 160	78.230	79.588	63.729	39.994	64. 953
	HEMBA1006914	54. 548	39, 053	48.945	35.736	25.895	38.586	22,479	33, 810
					29.982	32.625	61.537	62.750	30.818
30	HEMBA1006916	62.872	0.000	65. 115					
	HEMBA1006921	64.867	21.840	74.902	15.692	30.866	41.257	25. 569	10.362
	HEMBA1006926	51. 195	10.616	76.671	24.435	20. 300	84. 402	29. 503	20.967
	HEMBA1006927	24. 016	13.778	23.573	5. 335	15. 250	11.291	11.672	7.086
							5.970	5.688	3, 134
	HEMBA1006929	7. 146	8.487	5.431	5. 526	1.676			
	HEMBA 1006936	68. 233	22.847	45. 566	20.391	16.346	25.493	20.196	17.720
	HEMBA1006938	14. 202	8, 409	31.234	7.743	5.002	6.780	6.773	5. 945
35	HEMBA1006941		24. 290	40.928	13.779	16.040	34. 253	22.542	18, 507
	INCHEAT TOURS 41	30.559							65.640
	HEMBA1006942	147.487	57.842	121.883	69. 207	55. 456	76.853	61.942	
	HEMBA1006945	80. 546	64.930	104.037	63.709	40. 444	54.676	33. 533	31.915
	HEMBA1006949	10. 292	41.467	23, 921	1.860	15.813	7. 071	10.866	5. 231
	HEMBA1006952			34.750	8.032	18. 283	39.764	15. 332	12.456
		58. 685	12.572						34.791
	HEMBA1006960	91.939	38. 895	93.164	24.834	34. 400	36.160	36.715	
40	HEMBA1006973	74. 208	24. 793	50.621	17.619	22.844	24. 971	24.844	16. 167
	HEBBA1006974	48. 691	39,013	59, 414	48.064	16.799	38.579	21.301	45.006
	HEMBA1006976	35. 907	15.675	32, 116	19.091	14. 522	30.574	25.042	18. 348
								2.557	3. 536
	HEMBA1006989	6. 422	2.207	2.374	3.336	2.670	3.696		
	HEMBA 1006 993	334, 266	64.150	357.947	46.138	95. 466	144.777	109.174	54.000
	HEMBA1006996	9. 183	9.870	15.032	9.483	5, 722	9, 518	8.368	9.637
				224 000			47.863	27.205	56.828
45	HEMBA1007001	117.610	95, 668	334.868	56.093	55. 288			
	HEMBA 1007002	93. 134	41.846	72, 311	21.453	16.249	59.722	46.434	40.628
	HEMBA 1007013	65.734	23.106	53.712	16.933	20.783	34. 293	29. 163	29. 338
					6.385	9. 597	17. 982	16.658	15.035
•	HEMBA1007016	36. 649	14.972	27.491					
	HEM8A1007017	6.290	0.000	8.194	2. 155	5. 231	2. 329	1.949	0.000
	HEMBA 1007018	19.457	15.664	19.767	14.280	10. 586	15.084	9.105	14. 124
					15.913	53. 729	123.367	90.838	36.173
50	HEMBA1007044	139.784	50.078						
	HEMBA1007045	49. 576	7.913	39.757	9.069	10. 104	19.099	12.683	7.276
	HEMBA 1007051	36. 374	44.117	129. 384	27.586	19.407	24.088	15. 546	9.363
	HEMBA1007052	69. 582	19.611	40.507	19.050	9.213	19.409	18.969	10, 939
	HEM8A1007053	25. 326	27.611	21.861	14.031	14. 266	20.128	7.847	9.544
	HEMBA1007057	45, 897	13.545	33.857	18.616	25. 861	36. 241	14.769	13.902

Table 39

			. •		OIC 37				
	HEL'8A1007062	129.012	18.903	40.670	21.323	29.469	40.252	29.408	15,700
	HEMBA1007063	81.681	45.884	187.380	52. 391	36.943	28.608	35.303	41.236
	HEMBA1007066	98. 396	32.970	35. 373	22.961	11.085	42.430	26.631	14.760
5	HEMBA1007069	23. 449	21.519	78.409	16.835	27. 425	17.217	9.095	16, 163
	HEMBA1007073	54. 833	42.548	40.682	29.352	11.879	7, 937	24. 282	19. 372
	HEMBA1007076	83. 020	48.746	248. 260	61.189	50. 193	68.045	43. 836	
	HEMBA1007078	151, 561	159.600	445.445	189, 146	130. 283	98.734		35.650
	HEMBA1007080	43, 963	44.765	174.545				65. 934	117.079
	HEMBA1007084	78. 948			66.950	45.879	43. 194	43. 909	50.100
10	HEMBA1007085	263, 538	60.672	268. 327	63.769	63.088	60. 307	35.006	46.866
	HEMBA1007087		108.018	162.599	48.155	77.545	161. 321	63.614	80.640
		85, 598	25.085	47.862	25.580	13.918	62.815	143. 461	30.856
	HEMBA1007089 HEMBA1007095	21, 131	32.023	21.145	14.738	7.213	19.681	9.036	10.026
	HEMBA1007101	147.777	215.051	136.910	63.992	170,706	117.992	103. 152	86.452
		78. 959	53.790	147.891	35.676	28.082	27. 200	19. 131	25. 922
15	HEMBA1007104	66. 308	23.279	45.417	11.902	19. 468	48.054	26.760	16.647
	HEMBA1007106	28. 449	17.761	41.268	28.670	17.681	14.174	10.999	7.534
	HEMBA1007112	12.759	8. 412	16.340	9.319	7.661	7. 304	13.296	6.622
•	HEMBA1007113	126.702	0.000	229.408	64.551	40. 242	39.032	13.319	26. 174
	HEMBA1007121	219.036	207.410	696.658	149.217	168.827	131.628	642.099	128.755
	HEMBA1007129	50. 726	42.510	63.847	31.663	26.417	24. 371	18.928	20.103
20	HEMBA1007147	111.299	117.722	312.811	79.949	67.395	74. 391	35. 758	54. 184
	HEPBA1007149	83. 453	6.442	19.831	7.332	11.043	9. 349	9.831	8.756
	HEMBA1007151	97, 211	33, 530	53.944	24. 544	18. 501	35. 246	36. 228	24. 174
	HEMBA1007172	52.683	25. 324	438.704	42, 182	28.599	38. 126	26. 167	25.770
	HEMBA1007174	52. 921	13.482	44.770	21.384	19.520	28.559	22. 332	20.471
	HELBA1007176	89. 919	24.768	53.414	32.841	44. 643	73.679	87.040	30.762
25	HEMBA1007178	93. 941	73.120	135. 427	34, 313	32.040	34.622	22.898	24.897
	KEMBA1007185	62. 558	18.807	36,824	15.490	20.528	37.568	22.260	12.783
	HEMBA1007186	70.967	31.546	59.038	21.059	21.332	35.648	42.864	11.346
	HEMBA1007194	53. 376	38.911	126.660	33.992	23.875	21.109	12. 122	23. 307
•	HEMBA1007200	74. 955	53.829	44.212	23.979	20. 225	32.762	55, 417	22. 176
	HEEBA1007203	87.803	26.807	41.357	14.648	9.791	23. 392	30.167	17. 274
30	HEMBA1007205	82. 800	73.675	225. 293	44. 461	28.674	37.091	14.673	34. 505
30	HENBA1007224	25. 614	40. 402	50.116	21.484	14. 920	22.548	13. 197	20.053
	HEMBA1007226	88. 512	43.606	93. 121	22. 209	17.911	38.704	43.759	31.721
	HEMBA1007240	131.657	62.804	86.650	9.510	21.890	53.116	42.250	16.655
	KEMBA1007241	12. 225	7.719	18.461	5.051	6.724	15. 945	3. 135	5. 390
	HEMBA1007242 HEMBA1007243	21. 409	14.030	13.648	11.068	6. 265	17.370	8.487	5. 236
35		61.824	25.854	40. 264	17.235	23. 438	39. 197	31.904	20. 347
	HEMBA1007251	37.660	16.946	37.149	16.699	12. 180	19. 482	30.321	10. 262
	HEMBA1007267	53. 905	43.642	113.110	31.642	27.946	30.492	18.548	23.645
	HEMBA1007273	80.741	40.085	207.160	61.174	38. 220	29.008	32.292	29. 672
	HEMBA1007279	41.062 54.376	9.087	11.906	5. 193	6. 445	7.723	9. 225	4. 483
	HEMBA1007281		20.734	133.494	27.987	21.355	19.941	17.364	19. 503
40	HEMBA1007283	8. 523 25. 940	5.717	4.731	3. 403	2. 317	2.497	2.740	0.000
	HEMBA1007288	57. 959	14. 444 39. 576	24. 974 155, 227	23.487	19.771	23.418	19.378	26.409
	HEMBA1007291	37. 974	19.069	59, 253	28.725	24.689	25.110	16.998	16.095
	HEMBA1007299	446. 640	93.668	199. 852	61.423	13. 404 94. 129	17. 376 249. 345	13.060 241.373	13. 147
	HEE:8A1007300	103. 752	25.694	04 014				4.0	85. 323
	HEL'BA1007301	49. 752	18.178	32.677	18.217	33.650	25.018	31.407	10.569
45	HEMBA1007319	13. 312	10.598	23. 453	16.511	4. 278	33.786 9.382	22.892 2.996	12. 782 8. 570
	HEMBA1007320	53. 723	23. 595	62.301	29.439	16.672	32. 932	28. 191	
	HEMBA1007322	45. 986	125.362	77.545	43.693	17. 955		39.556	18. 418 80. 836
•	HEEBA1007323	64. 720	16.869	22.970			45.689 32.209		
	HEMBA1007326	313.094	189. 188		11.238 214.045	11.687		25. 350 70. 819	7. 506
	HELMA1007327	78. 767	61, 102	219. 980	55.002	178.109	171.587		115.174
50	HEMBA1007332	71.516	9.318			29.411	44. 095 24. 826	29.354	42. 286
	HEMBA1007341			34.879	5. 559	7. 452		12.763	20.050
	HEMBA1007342	89.805	53. 431 17. 289	207. 395	82.402	105.877	47.861	32.826	50.162
	HELBA1007347	22.063		28. 253	18.196	17.751	26. 378	13.820	9. 173
	HEMBA1007353	112. 392	64, 499	230.022	60. 348	47. 557	63.758	30.683	33. 285
		1.685	3. 520	0.575	1.850	1.976	0.107	2.061	0.788
EE	HEMBB1000005	60.047	46.027	121.870	38. 241	20.699	18. 268	20.068	26.957
55	HEU881000008	97. 929	53,604	274. 179	68.681	38.935	39, 328	26.881	34.873

Table 40

	DEUBBIOARATA I	122 120 1	107 001	120 100 1	120 410 1	57.867	95.203	75.902	92. 924
	HEMBB1000018	122.130	127. 851		120.419				
	HEMBB1000024	181.606	97.019	373.954	102.401	70.406	70.591	40. 304	66.798
5	HEEB81000025	85.919	29.049	45. 055	23.789	13. 946	24. 397	29.349	13.072
J	HEMBB1000030	108, 167	68.316	303.677	83,010	68. 378	81. 687	34.886	37.617
	HEBB100036	107.960	11.573	50.484	11.277	20. 480	41. 381	25.378	14.730
	HEMBB1000037	77.688	29, 380	69.658	56. 679	27. 020	54.062	30.086	15. 311
	HEMB81000039	52. 550	48.503	140.795	30.096	18.739	26.012	15, 151	21.723
	HEMB81000044	134, 136	75, 469	218.667	61.596	32.667	29.659	43. 360	42.831
	HELBB1000048	17. 937	21.052	31.004	18. 291	11. 321	20. 120	21.506	15.078
10	HEMB81000050	74.210	33.681	207.484	35.691	22.905	25. 584	18. 572	17.494
	HEMBB1000054	68. 273	47.191	248.350	44.008	24. 522	29.259	22.570	21.316
	HEMB81000055	72.875	112.284		110. 297	21. 358	70.636	93.824	132. 288
	HEMBB1000059	331.577	184.687	662.540	182.481	130.065	131.364	90.002	121.903
	HEMBB1000072	240.733	98.890	326.893	75. 919	67.742	118.222	108, 108	91.458
	HEMBB1000081	23.738	27.174	85.100	21.146	30.856	20. 458	7.513	15. 351
15	HEMBB1000083					37.789	59. 334	33, 712	39. 101
		120.759	58. 163	188. 224	40.609				
	HEMBB1000089	67.618	54. 952	191.832	56.629	24.609	36.847	30.680	26. 912
	HEMBB1000094	355. 534	116.828	161.958	31.504	29. 300	49.613	36.239	35. 197
	HEMBB1000097	27.834	63.724	51.488	14. 249	22.834	34, 068	18, 547	16. 455
	HEMBB1000099	157.641	91.912	456.470	71.078	50.739	64, 471	32, 108	43.354
	HEL:881000103	75. 781	59. 392	114. 974	44. 216	31.915	47.628	23.669	56.268
20									
	HEL'881000106	62.814	44. 996	77.918	35.044	19.825	40. 409	26. 156	45.001
	HEMBB1000113	43.660	33. 435	95. 987	42.744	19.714	20.114	15.899	21.606
	HEMB81000119	57. 350	21.211	42. 528	17.770	19.517	28.754	23.570	30. 104
	HEMBB1000133	92.950	65.230	58.619	69.544	53.706	104. 229	39.058	80.858
	HEMBB1000134	44. 120	20.654	76.693	40.611	24.712	37.185	42.327	21.963
	HEMB81000136	21.810	7. 191	44. 517	15. 599	7.339	22.582	12.399	24.899
25	KEMBB 1000141							36.737	52.602
		163.867	99.946	331.822	95. 807	55.858	64.560	30.131	
	HEMB81000144	96.831	97.019	183. 423	88. 529	36. 185	15. 577	29. 259	32.144
	HEM381000147	59. 253	9.088	62.426	7.391	11.451	7. 175	11.502	10.693
	HEMB81000152	56.391	28.723	34. 597	15.309	19.424	32.469	29. 105	19.117
	HEPB81000154	85, 308	47.878	101.061	33.881	19.477	27.298	20, 174	15. 366
	HEN881000155	35.691	36.132	109.038	28. 164	29.608	22. 283	16.557	17, 041
30									76. 388
••	HEMBB1000173	170.611	173.001		143.666	83.705	123. 932	65.317	
	HEE881000175	32. 273	19.114	23. 481	10. 948	4.039	29.180	7.135	13.322
	HEM881000176	56. 984	51.334	90.749	69.004	40.144	52. 980	25.845	19.359
	KEMBB1000198	70.426	12.768	26.381	10.237	6. 266	11.215	8.858	5. 363
	HEMB81000208	42.474	8. 955	34. 929	10.418	12.883	9.285	12.335	7. 978
	HEMB81000209	43.846	10.700	9.943	10.934	8.858	12.135	9.049	4, 168
35	HEMB81000212		12.579		15.361	33. 518	17.471	13.132	16.552
		27.532		76.077					
	HEMBB1000215	178. 324	89.053	294.606	95. 420	68. 598	89.720	51.270	61.235
	HEMBB1000217	148.073	45.416	96.614	47.569	37.572	89.989	48.073	33.510
	HEMB81000218	88.298	123.000	347.859	84. 124	41.828	57.417	21.147	34.605
	HEMBB1000225	70.693	14.949	41.586	31.786	30. 261	28.577	14.779	27.177
	HE::BB1000230	28.681	8.910	13.549	5.500	3.547	9.616	6.632	3. 293
40	HEM881000240	44.662	12.588	13.211	10.455	4. 589	41.554	8. 171	7.082
	HEMBB1000244	22.390	13.510	42.662	18.503	18.758	11, 192	2.111	13.188
									2.708
	HEMB81000250	20.878	6.254	20.741	9. 109	1.841	13.561	9, 540	
	HEY381000258	101.717	75.034	336.781	79. 281	52.303	67. 231	33, 313	34.880
	HET381000254	99.327	57.280	269.540	83.791	39.799	96.654	62.346	79.783
	HEMB81000256	70.747	23.082	23.217	14. 456	28.745	34.547	15.022	15.672
45	HEMBB1000272	14.990	14. 502	10.270	6.954	12.730	6.133	4. 205	16.611
40					49.759	41.568	43. 127	18. 199	25. 826
	HEMB81000274	105. 245	46.925	190.978					
	HEMBB1000276	6.479	2.218	2.501	4.783	1.754	2.070	2.079	1.252
	HEM881000284	4.790	5.088	7.884	3.489	2.213	3. 213	1.981	3. 304
	HEMB81000307	52.330	30.191	128.450	28.961	22.039	15.869	9.113	21.677
	HEMBB1000309	86.347	36.463	96.140	43.964	34.442	33.118	18.805	21.507
F0	HEMB81000312	41.862	30.986	40. 349	24. 933	7.383	79.360	24, 114	15.788
50									
	HEM981000317	49.311	18.053	26. 189	10.490	10.102	21.107	12.632	13. 384
	HEC:881000318	87. 180	33.847	208.954	43.556	23.043	27.764	9.191	17.641
	HEM881000332	3.892	11.256	14.087	42. 331	28. 145	14. 132	2.408	14.319
	HEM8B1000335	27.939	30.864	21.167	28.071	12.651	30.027	12.745	21.753
	HEM881000136	68.463	26.023	48.843	10.608	22.871	23.654	23.868	13.927
									60.614
55	HEMBB1000337	289.853	59. 290	93. 527	52.168	54. 197	125.769	126.562	00.014
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Table 41

	HEM881000338	£4 coc	75 755 T	122 400	44 E12 I	17.722	26.663	17,708	29.721
	HEM881000338	54.685 144.258	45.765 108.124	123.480	44.612 105.421	89.798	89.055	56.944	50. 241
	HEE:881000341				32.751	40.166	37.986	28.017	30. 881
5	HEMBB1000341	113.271	46.622	132.906		46.681	45, 761	43. 928	46.721
	HEMBB1000354	130.737	71.935	259.845 495.642	80.183 157.908	153.529	142.579	67. 161	105. 322
	HEMB81000358	202.146	151.264	29. 160		22.387	48. 989	59. 506	28. 803
	HEMBB1000369	92.244	22.827	97.758	24.670	21. 576	23.750	17. 278	16, 569
		55.720	25.874			38.792	44, 185	31.504	45. 653
	HEMB81000373	52.572	59. 105	70.779	61.379		80.319	58.214	75. 906
10	HEMBB1000374	153.545	115. 183		108. 150	98.073 60.328	63.876	73.647	43. 202
	HEL'BB1000376	95. 394	132.554	369.986	146.818	10.381	22.638	16.842	8. 781
	HEX:881000383	37.023	35. 429	24.954	13.017	30.891	57.827	37. 484	11. 921
	HEMBB1000391	127.327	30.055	106.971	24. 962	4, 100	8.569	2.643	8. 889
	HEMBB1000402	35. 143 82. 616	10. 865 20. 485	22. 406 44. 946	8. 561 25. 430	13.012	19.024	7.725	18. 695
15	HEMBB1000404	18.903	12, 568	10.300	8. 593	9, 455	9, 301	2.672	7. 956
15	HEMBB1000407	19. 286	8. 572	18. 593	3. 281	2.599	13.454	2.473	3. 407
	HEL:381000420	95.847	66. 573	138. 307	54. 950	39, 330	56. 220	37.608	43.081
	HEMBB1000430	274.820	161. 981	153.601	40.874	406.081	489. 107	693.805	115.638
	HEMBB1000434	350. 936	139. 481 1	599.497	199. 198	125, 426	113.500	65.776	77.687
	HEMBB1000438	67.342	10. 187	25. 472	7.736	8.148	27.875	7.217	6.701
20	HEMBB1000441	84.086	98. 109	312.643	78.842	60. 934	76.141	46.589	35. 267
20	HEMBB1000447	76.519	88. 156	54. 883	26.628	31.157	24. 328	25.777	38.008
	HEMBB1000449	22.367	11. 282	25. 245	11.267	1.700	13.053	5. 731	8. 109
	KEMBB1000453	26.781	29.875	49.056	22.139	35.305	22.456	14.005	15. 902
	HEMBB1000455	37.937	43, 401	129.423	29. 222	40.584	24. 577	21.227	20.356
	REMBB1000472	146.390	61.195	235.753	80. 306.	44. 122	82.882	52.783	87.457
25	HEMBB1000480	138. 135	67.904	194.466	46. 367	41.944	60.409	34.897	40.785
	HEMB81000486	78.511	63.045	211.876	47.786	39.049	36.558	20.396	17.632
	REMB81000487	21.510	22.091	29.116	10.718	21.Q56	15.854	13.086	10.892
	HEMBB1000490	232.419	148.116	562.064	159.218	134.370	107.851	60. 295	110. 306
	HEM2B1000491	149.070	107. 169	349.100	81.342	44. 330	51.147	33.633	59. 342
	KET'881000492	18.194	21.930	19.080	9. 690	6.821	10.632	9. 805	5. 454
30	HEI 881000493	286.390	34.074	64.876	31.406	23.065	49.816	39. 824	39. 921
	HEM-881000510	133. 225	95. 239	380.177	165.002	101.728	72.504	64.646	83.048
	HEY381000516	137.574	35.610	61.963	35.305	10. 932	78.851	39.905 1.611	19.224
	HEM381000518 HEM381000523	8. 388	3. 267	26. 133	5.489 82.474	1.531	1.500 69.756	32.830	51. 127
	HEL:381000523	153.793 46.151	88.071 13.390	329.880 40.950	8.319	32.799	6. 126	10.689	8. 426
	HEM3B1000542	57.808	36.831	46. 332	20.306	19, 414	5. 489	13.314	22.747
35	HEM381000550	39.123	26.036	79.169	22.945	10. 597	23, 147	37.266	20.568
	HEP381000554	192.214	105. 635	349. 184	148.874	90.632	98. 169	55.377	100.995
	HEMBB1000556	100.759	22. 180	68. 289	37.737	35. 176	41. 190	47.163	40.726
	HEMB\$1000564	101.412	37.586	144. 386	37.463	27.344	59. 939	31.447	9. 452
	HEC'881000567	361.516	76.515	125. 177	66.960	83.698	221.216	145.840	54. 204
	HEMBB1000569	63.847	46.712	54. 356	18. 197	23.752	36.942	31.264	39.479
40	HEE 381000573	99.088	89. 487	373.557	76.985	76.236	63, 534	39.587	58.648
	HEY381000575	74.071	67.726	296. 427	63.469	37.530	42. 388	33.544	46.151
	HEMBB1000579	27.868	12.805	18.934	6.889	3.743	24.452	24. 367	23. 262
	HEE'881000585	30.826	34. 244	65.882	26.172	19.828	26.184	16.826	33. 888
	HEY-381000586	85. 397	75.643	187.543	99.762	48. 456	35. 430	28. 693	50. 228
45	HEM 881000589	135. 404	58.619	243.853	51. 181	36. 284	29. 883	21.561	27. 997
70	HEMB81000591	99. 580	60. 946	242.306	54.695	36.589	52.616	32, 332	33.066
	HEMB81000592	30, 320	18.740	34. 338	11.753	8.732 46.829	28. 305 61. 565	49. 545	12.164 66.588
	HEMB81000593	148, 639	68.816	255. 892	61.084	9, 199	21.841	16.487	29. 580
	HEM881000595 HEM881000590	27.140	21.001 31.891	29.869 85.011	21.272	13.772	21.958	13.576	26.747
	HEMB81000611	39.074 14.828	6.552	11.601	7, 498	7.461	15.614	9. 246	9. 161
50	HEMBRIOCOSTY	193. 986	137.945		127.725	87.855	84. 583	46. 273	77. 986
	HEMB81000623	65. 566	26.480	50.777	1 19. 193	18.923	40. 974	28.571	23.219
	HEMBB1000630	62.606	23.074	40.815	18.796	14, 186	31.973	21, 492	13.779
	HEMBB1000831	61.311	41. 283	27.586	23.498	24. 433	35.043	48.566	22.826
	HEMB1000632	58.747	55. 433	156.750	30.460	29.661	33.497	21.899	21.857
	HEM881000636	127.885	47.562	59.456	48.965	33.643	65.366	42.360	37.349
55	HEMBB1000537	817. 391	628.017	1645.738	524.605	482.307	443.855	191.753	265.704

Table 42

	HEMBB1000638	55.058	47. 453	95, 751	42. 262	25.684	15.056	22.121	28.829
	HEMBB1000642	179. 188			80. 865	42,468	81.296	37.696	52.009
	HEX.BB1000643	43, 411			18. 985	11.038	14. 245	8. 276	18.743
5	HEMBB1000649	27.852		137, 371	34. 816	24. 496	9, 967	11.881	22.322
	HEMBB1000652	84. 942			78. 131	42.090	36.343	22.852	31.597
		418.308	73. 377	56.858	57, 166	32, 733	57.424	38.897	44.477
	HEMBB1000655	16.253	13. 954	10, 766	20.817	6.796	13, 110	7.987	4. 458
	HEMBB1000868			14, 606	13. 788	25. 844	15.049	12.549	11, 202
		28. 587 239. 020	13. 435 122. 952		19.970	96.244	75.058	66. 812	88.267
10	HE18B1000671				14. 904	5.916	4.811	2. 141	12.812
	HEMBB1000673	11.633	5. 779	14. 629 23. 438	7.697	1.049	30. 246	7.774	7.063
	HEMBB1000679	16. 899	7. 357		28. 150	66.411	89.722	49, 173	67.832
	HEMB81000684	188. 240			5. 085	1, 158	3. 240	3.421	1.785
	HEMBB1000692	4. 978	9. 265	11.569 59.522	22. 326	25. 408	13.898	31, 488	20.706
	HEMBB1000693	63. 119 15. 560	40. 561 31. 798	122.757	36. 451	19. 928	11.568	2.839	10.179
15		22.553	13. 626	23.777	8. 621	11.683	41.509	10.019	7. 584
	HEL'381000706		77.864	245. 726	50. 833	51.093	50. 427	37. 955	51.357
		74.737 23.726	10.733	6.625	12. 298	6. 349	9. 891	2.142	14.350
	HEMBB1000714	24. 239	9. 575,	11, 437	13.761	12.596	17.372	8. 105	16.144
	HEL1881000726	86.971	84. 395	208. 396	65, 157	43.881	37. 441	22.020	39.067
	KEL'381000729	51.556	25. 288	140. 931	23.005	27.775	18.629	12.838	14. 902
20	HELBB1000738	39.002	38. 955	166.616	42.588	21.380	43, 330	7.181	21, 192
	HEL'BB1000749	115.917	94. 942		136. 454	54. 340	39. 253	32.933	49.141
	HEM881000763	47.835	25. 201	36. 488	16. 952	21.036	31.919	14.990	12.111
	HEM381000770	30.598	45. 410	167.003	32.786	26.482	25. 698	18. 186	24. 127
	HEMBB1000774	27. 168	21.690	33, 470	20.937	12.916	22. 598	8. 092	17.606
	KEL'BB1000777	246.286	57. 131	58, 743	31.851	40.345	119, 113	81.364	53.990
25	HELBB1000781	41.945	36,620	34, 149	24.543	23.561	16.383	14.371	20.775
	HEMB81000788	10,756	10.608	5. 481	6.429	2.950	5. 995	4.522	4. 589
	HEYB81000789	28.490	9.620	26. 151	16.088	11.640	16.477	7.916	7.672
	HEY381000790	74.318	56. 925	185. 959	63.749	33. 523	24. 232	24.414	28. 423
	HET 381000794	18.080	17.254	38. 876	24. 305	7.427	10.338	5. 445	9. 305
	KE1381000807	50.070	31.869	22.751	19.865	20. 934	27.002	18.350	27. 280
30	HEMS 1000809	334. 541	42.976	42. 300	26.454	9. 545	31. 526	31.677	44. 152
	HEY881000810	189.365	50.676	163. 325	33.349	38.994	74. 400	45.398	19. 262
	HEMB81000821	40.710	9.304	21.006	6.841	5. 422	15. 981	10.835	5.685
	HELBB1000822	8.726	3.570	3. 541	1.411	7.255	5.519	1.285	1.525
	HELBB1000826	68. 485	40. 348	201. 149	68.467	43.204	31.769	32.812	55. 367
	HEYBB1000827	50.671	34. 326	108. 391	32.945	15.076	25. 813	18.713	25. 457
35	HEMB81000831	38.060	20.466	29.131	12.368	19.990	20. 562	25. 373 25. 687	6. 415 35. 108
	HEMB81000835	59. 181	56. 345	127.358	58.150	44. 350	35. 831 38. 995	20.712	30. 526
	HEMB81000840	117.639	63. 375	340. 802	61.186	48. 924 28. 984	47.603	29.642	29. 431
	HEMBB1000848	98, 938	53. 024	210. 423	42.569 2.559	1.621	1.272	1.364	1.086
	HEMBB1000852	1.827	2.160	0.621	14. 921	12.912	17. 279	10, 179	14. 915
40	HEMBB1000857	16.897 25.634	16. 768 16. 531	19.951 8.162	8.209	14. 482	12, 749	92.823	10, 102
40	HEMBB1000867	106, 946	56. 331	264.748	50.278	36.949	41. 202	26. 795	29.760
	HELBB1000870	68. 550	62. 423	192.351	52,406	39.303	55.641	23.738	27.427
	HEMB81000875	21.813	12.044	24. 968	11.314	7.689	10.690	11. 143	26.241
	HEMBB 1000881	30.089	16. 478	28. 345	14,926	18.419	17.763	18, 901	20.494
	HEMBB 1000883	11.669	10.263	26. 185	6.975	2.780	8. 223	2.906	3.540
45	HEL BB 1000BUY	42.638	32. 274	66.780	22.979	31.512	42.842	20.622	22.566
40	KEMBB1000888	20.318	8. 193	11.483	5.178	4.073	8.708	6.801	4. 342
	KEL BB 1000890	40, 795	42. 287	112.076	25.031	11.171	23.116	15. 491	16.447
	KE 881000893	38. 227	10.603	88. 306	24.535	14.440	12.863	9.734	17.727
	KET'881000900	23.814	8,709	17.013	9.267	10.928	12.199	14. 105	11.108
	HEL'381000905	63. 589	43. 501	37.125	41.367	26, 379	29.649	38.699	31.891
50	HEY381000908	42.944	54. 674	120. 821	34.982	28.838	28. 194	15.897	26.230
-▼	HEL'BB1000910	72.960	51.795	161.850	41.050	36. 594	37. 378	13.612	23. 263
	HET'881000913	33.820	35. 219	96.448	24.688	12.371	26.067	14.715	19. 268
	HEP 381000915	1910.513	222. 511	693. 345	124.825	532.993	1548. 228	1159.943	223. 176
	HELB81000917	99. 638	64.212	310.142	53.316	39.091	34. 989	22.324	40.667
	HEL BB 1000927	80. 569	11.252	19,448	8.653	21.944	24.546	17.769	17.391
55	HEMB81000932	33. 128	33.556	95.029	29.041	17.945	21.758	22.973	31.034
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Table 43

	HEMBB1000933	883.639	393.035	605.052	289. 543	312.660	538.431	353.155	291.706
	HEMBB1000936	23. 212	17. 243	46.380	14. 205	25.527	13.908	8.530	11.716
5	HEMBB1000939	105.016	36.905	52. 525	19.304	30.111	35. 223	41.856	37.146
3	HEMB81000941	6.540	27.555	15.872	4.660	6.130	17.648	83. 246	9. 541
	HEMB81000947	36. 384	18.020	47.143	21.361	9.585	34. 299	13.482	13. 269
	HEMB81000954	16. 970	17.878	19.810	11.407	6.851	17. 302	10.023	8. 877
	HEMB81000959	22.223	21.226	78.296	22.443	5.599	10.268	10.673	12. 183
	HEMB81000973	11.584	10.364	21.189	8. 579	7.102	23.845	5. 510	9.891
10	HEL 881000975	99.598	37.022	69.027	23.084	27. 137	40. 162	56.997	30.316
	HELBB1000981	10. 199	12. 524	23.602	20. 141	5.813	6. 152	13.771	4. 102
	HEMBB1000985	13.065	8.026	7.574	4.776	6.642	2, 985	6.049	3.612
	HEL:881000991	67.124	17.092	28.053	8.864	8. 560	28. 394	25.072	10. 907
1	HEMB81000996	170.256	127.636	352.650	90. 350	64. 926	71.240	7.698	102. 622
	HELBB1001000	48. 257	19.380	16.573	15. 226	10.611	14. 541		9.642 0.000
15	HEY881001004	0.797	1.839	0.439	0.000	0.000	0.318 12.238	0.000 6.478	9. 235
	HEMBB1001008	17.533	13.975	16.434	11. 194	6.400 15.302	17.720	15. 586	16.702
	HEMBB1001011	39.743	19.337	28. 395	15. 752 50. 619	33.004	55.708	30, 100	34. 469
	HEMB81001014 HEMB81001020	121.726 86.065	46.352 68.022	244. 715 243. 352	67.763	53. 522	50.406	30. 247	49. 844
	HEUBB1001024	66.546	59.010	205. 347	41. 480	31.865	35. 052	21.045	39. 489
	HEMBB1001026	36. 265	27.027	76. 443	19.990	25. 484	27.657	12.014	23. 129
20	HEMBB1001037	64. 392	37.810	120.090	20.652	22.459	27. 294	18, 918	28.917
	HEMBB1001042	58. 936	20.428	42.468	17. 255	15.600	32.463	20. 274	18.506
	HEL'881001046	76,790	22.021	40.791	13.932	17.825	47.853	26.672	30.056
	HEMB81001047	76.665	39. 237	208.757	53.469	44. 539	37.624	16.049	20. 262
	HEMBB1001048	133.028	58. 176	140.515	48.390	34.614	42.111	29. 526	34. 858
25	HEMB81001051	22.699	8.465	13.142	9. 942	10.065	9.946	5.881	8. 790
20	HEMBB1001056	40. 040	16. 494	45.000	22.674	18.685	21.131	18. 431	12. 498
	HEMB81001058	88.873	59.116	223.822	45. 122	34.696	29. 783	21.562	25. 222
	HEMBB1001060	35. 486	18.631	33.852	60.851	26.807	13.499	12.993	19.391
	KEMBB1001063	53.418	36. 359	125. 166	33. 156	24. 220 26. 835	19.182 79.006	63. 198	43. 296
	HEMBB1001068	79.181	46.879	78.756 173.393	35.034 49.675	25. 253	33.015	14. 189	22.904
30	HEMBB 100 1082	66.296	58. 491	20.825	17.116	14. 939	41.581	21. 497	13.792
	HEMB81001095 HEMB81001096	64.435	31.409 28.562	34. 366	32.120	13.089	21.236	15.814	22.034
	HEMB81001101	79.652	21.131	40.775	18.757	35.350	46.263	18. 855	13.874
	HEMBB1001102	51.740	27. 685	86.794	21.160	12.958	16.450	7. 235	8.605
	HEMB81001104	61.846	33.489	28. 997	14.789	10.623	20.859	15. 993	10.658
	HEMBB1001105	69.199	32.868	132.855	27.292	32.605	49. 984	20.779	23.761
35	HEMBB1001112	161.356	78.361	73.588	64.617	86. 150	93. 363	87.696	95.854
	HEMB81001113	114,744	130.208	298. 139	107.218	73.757	61.718	32.824	66.952
	HEMBB1001114	105. 358	95.960	365.719	66.457	62.314	35. 251	34. 480	51.970
	HEM381001115	67.274	16.815	13. 190	26.838	17.638	29. 948	23.803	34. 239 18. 952
	HEM881001117	2.434	10.619	14. 951	4. 152	4. 937	2. 694	2.729 5.261	9.614
40	HEMBB1001119	18. 198	17. 501	58.077	15.560 81.302	5. 202 76. 905	13. 437	58.863	61.487
40	HEMB81001126	306.301 39.673	111.345 36.703	266.365 178:312	45.328	36, 363	38.712	14. 400	26. 997
	HEMBB1001137	53. 424	19. 209	46.849	14. 453	13.705	30. 395	18.865	15.761
	HEMBB1001142	105. 888	131.411	405, 403	98.008	104, 700	62.754	12.598	75. 485
	HEM881001145	114.864	106.329	348.161	78.364	57.587	54. 983	24.738	51.568
	HEM881001151	149.618	23.632	66.607	14.582	34. 238	68.060	46.084	19.806
45	HEMBB1001153	92. 263	53. 444	153.351	44.131	37. 191	34. 991	21.708	32. 599
	HEMB81001158	64.416	30.844	50.578	22.880	32.523	47.046	24.553	39.658
	HEMBB1001169	95. 424	70.158	253.814	76.490	44.058	37, 113	24. 102	38.757
	HEMB81001170	34. 989	7.730	32.617	5. 324	4.217	11.418	7.623	5. 208
	HE1281001175	46.512	27.401	45. 252	21.001	15.416	20. 636	17.361	36.021
	HEMB81001177	126. 389	85.212	396.633	84. 357	48. 470	40.910	34. 438 25. 771	42.680 26.488
50	HEMB81001182	70.825	30.508	45.077	19. 262	28.316	32. 507 22. 456	24. 299	31.214
	HEY-8B1001192	30.059	21.703	61.610	20.151	5.688 0.797	2. 148		1. 223
	HEMBB1001199	1.469	0.000	0.000	4.430 5.734	0.000	2.413	1.567	2.969
	HEM381001200 HEM381001208	2.266	1.426 37.738	122, 154	28.426	28.653	55. 253		21.624
	HEMBB1001208	103.602	77.445	233.649	60.849		40. 993		
55	HEPBB1001210	14. 499	40. 527	32.902	6.231	10. 125	16.413		28.930
55	11000001001210	1 7. 700	1 .0. 021	1				-	

Table 44

	HEBBB1001215	219. 922	83.033	126.326	63.007	71.733	115.441	61.961	72. 230
	HEMB81001217	63.633	22.116	41.047	17.479	20.160	53.164	31.645	18.739
5	HEMBB1001218	98. 226	47.137	142.256	53.412	29.467	23.819	20. 495	24.079
· ·	HEMBB1001221	0. 524	1.310	12.795	0.988	0.992	0.867	0.000	1.767
	HEMBB1001224	52, 109	37. 281	86.318	28.364	24. 177	19.072	16. 478	20. 321
	HEMBB1001230	38. 785	17.158	30.714	15.256	12.698	31.469	27. 596	17. 436
	HEMBB1001234	335.966	64.817	131.669	43.601	69.385	167.134	101.415	57. 258
	HEMBB1001235	152.870	67.952	84.726	40.262	26.665	52.686	38. 623 18. 264	49.693
10	HEMBB1001237	16. 971	23, 623	33.663	30.744	21. 161	18. 495		7 500
	HEMBB1001242	26. 787	15.776	22. 922	4. 200	5. 187	11. 277 4. 366	2.735	7.589
	HEYB81001244	280. 439	9.589	9.743	8. 128 25. 983	2.116 19.890	21.254	15. 839	21.542
	HEM881001249	51.892	27. 766	106.010	31.656	8. 253	38. 144	20. 639	25. 942
	HEMBB1001253	50, 869 28, 109	33.773 8.716	58.857 61.080	12.779	6. 376	18.461	22. 558	8. 559
	HEMBB1001254	2.010	9.088	3. 704	1.682	16. 420	18.653	1.717	1.611
15	HEM881001266 HEM881001267	131.334	93.697	391.730	88. 886	45.610	62.418	33.457	63.350
	HEMBB1001271	31, 480	28. 408	63.773	19.821	15. 244	12.530	8. 683	10.739
	HEMBB1001282	41, 166	11.440	25.546	10.847	7.531	21.762	15, 737	10.592
	HEE881001287	195. 274	200.678	131.870	63.454	15.491	70.758	43.360	52.931
	HEMB81001288	40. 232	10.227	25.481	9.789	5. 520	21.519	16. 538	9.861
	HEMBB1001289	84.233	74.730	246.417	61.615	31.689	36.447	24. 521	38.077
20	HEMBB1001290	57.742	13.181	11.174	33.921	23.320	24.860	82.615	15. 369
	HEMB81001294	80.761	23.745	72.937	16.689	20. 147	45. 268	37. 586	22. 951
	HEL'881001299	58.616	17.094	44. 424	13.532	14.650	31.325	32.822	12. 329
	HEMB81001302	87.107	24. 979	56.357	23. 389	20.784	37. 921	28.849	21.981
	HEM881001304	12.134	0.119	5. 246	19.403	1.810	3.978	2, 153	1.580
25	HEMBB1001314	6, 410	5.111	25.042	5. 961	3. 244 1. 405	7.037	2. 954 1. 659	2. 258 1. 943
	HEMBB1001315	3.706	8.398	10.733 87.084	3.067 32.290	25. 473	21.551	14.009	18.118
	HEMB81001317	13.902	34.918 5.726	7.704	2.886	2. 324	1. 546	2,008	5.612
	HEMBB1001331	34.871	17.866	37.859	11.626	6. 188	23.138	24. 975	17.786
	HEM381001335	22, 550	20, 911	19. 341	12, 458	15.964	18.477	15. 941	5.614
	HEM3B1001337	61.645	43.894	187.675	45. 250	52.185	20. 178	25.750	29. 233
30	HEMBB1001339	20.634	25.030	21.230	11.541	12.874	18. 490	12.601	13.466
	HEMBB1001344	31.209	8.322	15.710	5.412	6.749	16.517	16. 482	9.869
	HEMBB1001346	44.149	21.512	38, 191	15.415	9.432	26. 936	17,706	15. 965
	HEMB81001348	66.624	40.319	173.356	39.887	26.835	31.783	20.641	26.670
	HEM8B1001350	103.603	17.400	35.832	13.555	13.837	54. 503	34.694	19. 925
25	HEMB81001356	12.440	11.385	25. 095	8. 592	6.787	7.806 12.620	8.759 13.025	8. 923 10. 117
35	HE1381001364	28. 525	14.483	31. 452	11.829 52.112	13.494	41. 191	29. 156	32.064
	HEMBB1001366 HEMBB1001367	57.883 140.660	53.690 59.744	210.263 283.101	54. 260	46.338	67. 368	43, 944	48.485
	KEMB81001369	17. 341	20.708	71.044	14.855	7.629	12.537	7.158	14.407
	HE1881001380	50. 204	67.647	124. 463	41.290	43.730	41.591	29.026	63.358
	HE3881001381	19.588	19.545	34.218	14.113	18.710	9. 428	10. 202	13.801
40	HEMBB1001384	17,779	11.154	26.926	11.606	19.030	10.038	7.367	14. 535
•	HEMBB1001387	20.705	16.837	19.148	9.955	8.901	15. 994	7.831	13.345
	HEMBB1001394	21.419	19.091	32.720	17.551	19.172	11.590	12.282	11.322
	HEMBB1001407	39, 158	17.718	75.721	24.299	17.481	17.410	20.342	15. 525
	HEI:881001410	18.880	3.346	6.042	2.907	2.655	0.000	2.839 12.610	2.094
	HEMBB1001413	32. 291	25.769	80.279		21.102	11.132	13.895	31.342
45	HEMBB1001419	36.323	42.415	185. 239	24.790 12.065	15.685	17. 972 64. 181	165. 647	23. 322
	HE1381001421 HE1381001424	29. 464	57. 495	109.370	6.073	6.773	7. 183	5. 215	8. 524
	HEMB81001426	9. 663 36. 471	7.148	86.872	20.138	17.823	19.534	15. 347	23.782
	HEMB81001429	60.351	47.669	39. 928	29.802	21.695	39. 456	39.474	41.210
	HEMBB1001436	168. 445	86.814	350. 902	88.825	54.546	86.724	48.813	58.527
50	HEMB81001443	20, 733	11.137	12.445	8.769	16.707	14.531	9.581	12.477
50	HEM8B1001449	70. 239	34.064	146.511	28.311	23.391	19.979	16.080	22.377
	HEMBB1001454	60.851	40.766	133.878	33.168	28.709		29.720	26.623
	HEMBB1001458	77. 938	28.808	33.472	15.970	29.260	40.965	25. 268	28.079
	HEMB81001461	44. 192	44.580	179.531	65.974	16.217			27.974
	HEMBB1001463	57. 949	102.937	230.980	60.751	41.957		25. 233	38.517
55	HEM881001464	18.058	9. 999	14. 908	10.039	7.528	8.680	2. 638	2.964

Table 45

	HEMBB1001456	31.340	22.324	20. 480	15.496	3.611	15.533	10.020	13.761
						4. 887	24.039	4, 114	4, 470
	HEMBB1001482	12.741	4.057	9. 987	4. 175				
5	HEMBB1001500	26.823	21.417	65. 107	17.492	9. 196	12. 958	6. 167	14.603
	HEMBB1001505	116.783	105. 297	302.199	104. 682	36.419	54. 346	38. 027	46.591
	HEMBB1001521	55, 379	38.602	133, 188	25.792	20. 204	23. 504	18.528	22.786
	HEM8B1001527	331.186	160.160		131, 308	116.694	179.333	72.732	79.869
	HEMB81001530	24.722	25.693	57, 090	19. 457	7.662	20.875	31.031	23.503
					34. 674	21.061	27.704	18.966	32.578
	HEMB81001531	43.913	51.679	130. 225					
10	HEMBB1001532	6. 957	3.901	34. 322	7. 593	1.875	8. 172	300.808	7.501
	HEMBB1001535	71.654	59. 202	131.794	46.369	28. 936	34.644	21.690	23.017
	HEMB81001536	73.109	48.204	106.813	35. 175	16.411	22.356	19.126	20.785
	HEMBB1001537	40. 809	54.756	140.043	43.830	21. 583	31.273	8.692	29.500
	HEMBB1001542	79. 436	33.152	94. 294	34. 360	26. 100	44.300	19.679	22.657
	HEMBB1001543	55. 819	14. 588	8.417	4. 239	7, 702	20.740	11.834	18.032
				12. 415	9. 202	10, 101	15.047	10,631	8. 198
15	HEMBB1001547	10.746	8.433				115. 789	76, 174	67.211
	HEMB81001548	163.125	42. 223	39. 134	33.781	26. 421			
	HEMB81001551	32.248	10.176	8. 937	9.728	20.037		7078.074	11. 439
	HEMBB1001555	62.998	58.959	166.842	57.865	40.731	30. 981	17.189	40.721
	HEMB81001562	67.088	35. 544?	83. 929	24. 475	18. 852	28.472	27.682	23. 295
	HEMBB1001564	139.467	320. 422	580, 390	304.052	124.857	300.720	202. 502	439. 361
••	HEMBB1001565	56.749	43. 545	123.727	39, 891	29.530	30.029	17.527	28. 501
20	HEMB81001569	34. 482	26. 904	100.487	28.883	16.462	19.020	8.403	16.605
	HEMBB1001573	48. 940	40. 308	65. 598	41.979	32. 247	35. 238	25.583	36.979
			57.831		61.076	40. 832	38, 446	18.915	42.636
	HEMBB1001585	153.364		211.685			24.673	16.535	26.124
	HEMBB1001586	44.946	40. 343	113. 224	34. 426	18. 386			75. 499
	HEMBB1001588	157.947	130.811	402.650	111.293	69, 831	80. 240	46.050	
25	HEMBB1001595	12.602	11. 160	44. 464	13.949	6.811	11.538	4. 359	11.569
	HEMBB1001596	53.986	20.798	39.629	25. 473	20. 578	32.621	23.309	36.564
	HEMBB1001599	29.275	7. 352	13. 267	11.568	5. 279	15.756	10.260	5.135
	HEMBB1001603	3.581	2.642	7.782	4. 279	3.051	0.341	1.424	3.160
	HEMBB1001606	6.897	7. 220	7.226	7.657	3.104	5. 383	5.658	4.364
	HEMB01001612	101.576	58. 128	240.469	58.770	36.287	42.917	27.221	40.063
	HEMBB1001618	52.604	38. 648	141.745	37.723	24.274	24.922	17, 197	24.223
30	HEMB81001619	59. 431	78. 268	138. 545	63.285	52, 275	37.035	22.185	38.081
	HEMB81001623	33. 128	8. 489	11. 122	6.318	8. 326	16.007	3.331	7.918
					7.577	2. 293	8. 389	1.716	4.647
	HEMB81001625	10.068	16.076	8.496					5.942
	HEMB81001630	7. 144	5. 464	31.186	8.383	3. 256	11.196	3.053	8.859
	HEM881001635	18. 151	8. 186	33.138	13.501	9, 143	9.688	44.037	
0.5	HEMB1001637	40.224	35, 174	58.964	24.082	26.640	26.340	20,792	26.243
35	HEMBB1001641	21.655	10.768	33.553	9. 122	5.845	7.210	5.796	8.300
	HEMBB1001653	76.468	45. 984	138.114	33.606	30.023	33. 136	16.720	25. 949
	HEMB81001665	3.000	0.352	5.654	0. 275	0.718	0.106	0.899	0.407
	REMBB1001666	48.027	23. 276	59.669	22.201	9.196	20.512	10.659	15.687
	HEMB81001667	2.570	7. 909	3, 107	5.847	8.690	2.748	1.999	8.738
	HEMB81001668	2.545	8.886	13.392	8.498	18, 131	3. 355	1.531	3.932
40	HEMBB1001669	5.751	5. 364	10. 395	3.219	4. 970	5.110	4.341	2.139
	HEMBB1001670	17.795	10. 903	34, 891	20.715	11.725	22.401	12,909	20.514
	HEMB1001673	69.924	44, 194	58.806	53,036	21.640	40.433	25.038	49.339
						9. 356	22.270	15.894	11.977
	HE'881001675	58.961	13.650	21.648	10.914	11, 101	24.642	13.266	4. 383
	HEMB81001679	51.245	9, 166	29.461	6.718				13.683
	HEE 881001684	27.854	11.218		14.666	11.546	25. 422	15.072	
45	HEM381001685	9.626	8. 721	34.446	7, 134	4.659	1.316	3.180	6.172
	HEMB81001695	2.706	4. 723	4.741	1.162	8.059	1.109	1.036	1.119
	HEMB81001703	116.774	37.756	115.693	36.901	34.790	69.383	44. 901	43.576
	KEM981001704	67. 385	52.606	211.228	52.452	40.406	43.432	33.952	54.662
	KEMBB1001706	122. 282	70. 476	227.746	77.627	63.608	53.010	38.740	56.789
	HEM881001707	111.416	69.815	154. 286	51.656	60.773	50, 260	33.306	43.746
50	HEMBB1001717				10.609	5, 688	9. 921	4.816	8.073
50		14. 112	16. 260	60. 454			35.654	37. 976	32.089
	HEYB81001731	29.550	36. 222	21.992	33.872	22.551			17.320
	HEMBB1001734	75.818	39. 477	107.419	26.507	15. 856	20.715	17.010	
	HEMBB1001735	63. 245	22. 136	169.823	34. 289	26.478	18.371	17. 292	27.924
	HEYB81001736	20.722	18.061	27.944	17.598	12.534	9, 551	10.504	13.178
	HEL1881001747	21. 158	15. 281	18.501	9.967	9.806	11.088	17.268	12.572
55	HEMBB1001749	89. 421	90.342	429. 206	126.585	53.728	56.733	28.560	64.467

Table 46

	HEMB81001753	85. 135	63.020	101.881	44.766	60.100	46.138	48. 988	44. 990
	HEMB81001756	86.556	37.048	83. 531	33. 276	42.763	54. 273	32.005	30.821
5	HEMB81001757	1. 981	3.522	5. 232	3.590	1.394	7.486	3. 256	3.014
5	HEMB81001760	13.573	14.554	27.053	7. 204	5. 280	8. 129	5. 242	4.088
	HEMB81001762	26.210	15.945	24. 826	8. 457	6.461	26.934	6.893	9. 656
	HEMB81001780	18.738	33.363	27.562	17.311	13.893	4.277	14. 584	19. 429
	HEMB81001785	3.266	2.954	7.974	3. 522	3.900	7.429	3. 964	4.008
	HEMBB1001788	77.710	51.716	232.298	72.096	40. 555	41.418	29. 586	33. 423
10	HEMB81001793	221.348	29.215	45. 528	20.500	22.918	33.927	36.095	25. 245
,,	HEMB81001797	4.049	9.015	10.442	4.015	2. 532	8.773	2. 904	6. 333
	HEMB81001802	430. 563	24.213	34.832	14. 183	17. 392	26.448	23.001 15.772	29.744
	HEMBB1001812	91.804	71.389	218. 174	56.457	56.645	54.459		55. 255
	HEM881001815	506.853	426.652	275. 995	120.005	129. 468	289.852 45.331	148. 011 35. 194	122. 368 47. 899
	HEBBB1001816	90.698	55.478	178. 334	52.637	25. 170 9. 329	19. 975	9, 745	18.634
15	HEMBB1001831 HEMBB1001834	22.874 456.615	14.551 299.793	45. 474 406. 927	16.825 241.146	284. 283	499. 103	267. 485	306.611
	HEMBB1001834	138. 292	91.469	348. 309	101.544	73.058	67. 103	40. 539	76. 261
	HEMB81001839	9. 720	6.600	7.318	0.000	2.606	4. 296	2.217	2.738
	HEMBB1001841	345. 524	134.230:	67.049	25. 938	60.560	21.530	21. 177	18, 486
	HEMBB1001844	61.041	25.820	34.819	14.237	14.648	34.333	20.655	31. 102
	HEMBB1001847	126.241	111.341	239.722	147.873	65.849	85.164	47. 980	108. 378
20	HEMBB1001848	40.802	39.856	24.837	12.646	9.727	18.893	18.093	17. 754
	HEMB81001850	171, 151	101.141	118.680	33. 622	64.050	118.354	50. 599	75. 857
	HEMBB1001859	133.676	77.853	231, 163	65.024	41.660	123.173	103. 961	48. 695
	HEMB81001863	115.353	92.421	255. 141	83.601	85.833	53.693	30.832	49.888
	HEMBB1001867	15.427	15.822	8.336	10.061	4. 673	8.415	6. 299	9.816
25	HEMBB1001868	24.470	17.457	24. 238	7.996	8.810	8. 133	10. 520	11. 923
25	HEMBB1001869	82.894	76.711	234. 322	61.007	44.801	45. 547	29. 853 9. 542	39.008 5.964
	HEMBB1001872	15. 921	7.288	5.998 22.113	10. 151	2.561 9.515	5.674 14.138	6. 058	5. 891
	HEMBB1001874 HEMBB1001875	36.336 7.615	11.065	13.755	15. 221 26. 314	11.646	3.662	5. 863	7, 228
	HEM381001880	107.638	82.806	115.014	59. 163	39.712	47.440	27, 454	37.214
	HEMB81001899	15.785	11.630	15. 181	7. 571	2. 259	12.203	4. 190	3. 366
30	HEMB81001903	59.215	24.149	27. 564	15. 205	8.601	28.805	15.592	15.765
	HEMB81001905	29. 932	24.402	20. 256	15.117	8. 559	17.138	12.021	12.009
	HEMBB1001906	15. 456	13.077	51.260	10.147	16.547	10.906	7. 943	9. 129
	HEMBB1001908	35.095	32.316	100.465	26.514	24.742	20.649	8.759	14, 223
	HEMB81001910	67.419	35, 922	139. 126	58. 266	43. 100	26.178	19. 330	29.710
	HEE:381001911	50.456	46.682	196.311	58. 337	31.782	35.278	19.934	32.009
35	HEMB81001915	40.796	27.017	19. 351	20.885	15.345	12.662	9. 798	36.052
	HEMB81001921	95.398	115, 190	314, 157	85.049	59.940	59, 397	36.034	60.585
	HEMBB1001922	54. 587	37.299	107.814	29. 796	15.712	23.741 16.405	15.662	16.568
	HEMB81001925	35. 478 9. 272	39. 156 7. 467	106.631	7.045	3. 402	5.636	2. 969	5. 808
	HEMBB1001944	122. 259	83, 163	268. 572	86. 582	66. 995	51.236	27. 262	45. 542
40	HEMBB1001945	55. 555	20.668	28.702	7.169	21.076	24.208	18.042	10.472
40	HEMB81001947	47.254	12.987	21.887	16. 223	6.133	25.673	16.697	13.440
	HEMBB1001950	99.345	31.711	42.202	32.724	17.168	58.211	28.763	30. 429
	HEMBB1001952	67.117	40.169	164.691	39. 168	16. 287	31.103	11.276	24. 511
•	HEMBB1001953	56.049	47.572	147.635	34. 659	22.662	21.660	13.445	22. 280
	HEMB81001957	43.669		106.261	26. 369	15.837	16.589	5.199	12.837
45	HEYB81001959	25.731	45. 573	72.402	48. 003	21.477	24.564	17.194	36. 361
	HEMB81001962	59. 585	38.413	125.747	48. 471	52. 786	45.598	20.834 46.606	29. 320
	HEE'BB1001967	156. 252	96.306	460.639	121.361	89.090	70.066	24. 193	68. 839 43. 482
	HE13881001973	62.418	55.111	203.353	61.777	40.564	56.711	62.043	55. 171
	HEMB81001978	205. 611	67. 998 97. 908	184.804	55. 506 79. 417	69. 496	62.957	41.995	65. 291
	HEMB81001987	23.094	30.009	63.743	16.838	10.970	10.414	5. 543	10. 645
50	HEMBB1001988	26.549	17.876	71.399	12.651	11.631	11.873	6. 563	10. 248
	HEMB81001990	61.049	28.808	125.791	31.477	30.752	26.525	9. 894	24. 366
	HEMB81001996	40.435	12.303	17.096	14, 159	3.837	18.573	11.696	13. 433
	HEMB81001997	91.453	62.313	247.838	54.724	40.131	29. 522	27.492	42. 942
	HEYBB1001999	28. 583	9.839	33.748	34. 520	11.455	23.048	14, 798	25. 158
55	HEMBB1002002	19.354	10.115	14, 415	9.527	16.781	12.044	7.088	14.724
JJ									

Table 47

	HEMBB1002005	127. 202	87.407	314. 165	82.406	66.505	55. 577	40.792	64. 185
	HEMBB1002009	0.000	1.364	22.770	0.807	4. 369	1. 295	0.000	0.000
	HEMBB1002013	28. 258	13. 576	16.813	10.399	10.765	17.046		
5								7.782	9. 691
	HEMBB1002015	105. 576	48. 524	66.937	36.377	38. 220	74.637	28.221	34. 621
	HEMBB1002024	216.724	27.841	16.159	12.961	10.268	16.725	13.378	30. 580
	HEMB81002035	46. 139	20.267	93.090	25.830	19. 155	14. 290	9.089	10.861
	HEMB81002039	56.819	33.510	91.779	23.686	12.816	13. 451	13.710	16.666
	HEMBB1002041	64. 639	34. 426	51.061	22.611	27. 241	31.364	25, 209	28. 240
10	HEMBB1002042	108. 989	70.262	244.087	61.596	54.097	58. 195	45. 407	53.478
70	HEMBB1002043	45.022	36.752	179.777	48.242	21.779	25. 603	30.919	28. 446
	HEMBB1002044	13. 181	2.012	5.797	1.053	1. 982	1.313	3. 432	2.045
	HEMB81002045	289. 530	197.322	441.790	143.182	150.349	206.083	108. 290	118.515
	HEMBB1002049	35. 193	24.481	83.015	26.999	19.710	27. 535	16.278	24. 921
	HEM8B1002050	37.095	16.954	49, 110	12.868	13.580	16.690	9.422	14, 540
	HEMBB1002051	36.389	19.655	68.218	18.665	8,800	22. 352	16.403	17.616
15	HEMBB1002068	75. 935	30.174	53. 312	27.588	23.758	28. 553	40. 522	36.664
	HEMB81002069	213.038	176.212	471.114	127. 141	113.252	145.813	82.555	84. 929
	HEMBB1002075	42.631	31.316	161.071	28.782	21.239	25. 996	13. 087	18. 589
	HEL'881002079	16. 958	10.592	15.974	7.658	4. 913	11.054	12.406	9. 170
	HEMBB1002080	43.775	32.579	72.576	24.001	9.827	28.608	17.214	17. 433
	HEMBB1002082	26.775	8.257	21.193	4. 448	6. 280	19.090	464. 903	8. 346
20	HEMBB1002084	17, 127	6.840	43.925	4.043	9.757	26.316	9.627	6. 512
	HEMBB1002088	90.318	38.977	65. 816	40.755	47.974	81.367	57. 452	75. 281
	HEMB81002092	192, 949	59.522	268.965	49.978	47. 797	60.595	48. 524	38. 080
	HELTBB1002094	127.875	84.707	379.671	89.066	80.779	70.636	38. 807	57.037
	HE11881002103	29.830	9.307	18.867	12.419	117.011	11.825	10.555	6. 133
	HEMBB1002109	28.380	23.579	104. 568	24. 307	17.018	17.089	11,301	21.844
25	HEMB81002115	71.073	86.440	117. 523	95. 976	28. 307	85. 908	60, 445	114.378
	HEMBB1002120	16.393	10.090	4. 147	2.085	3. 568	9. 594	4. 954	4. 539
	HEMB81002121	12.050	2.757	6.522	1.146	2.007	0.000	1.999	1, 549
	HE2381002134	784. 781	365. 377	605. 805	262.168	223. 204	719.592	534.370	450. 949
	HEYB81002136	109. 220	32.405	75.010	27. 402	26. 278	36. 231	38. 283	23. 593
	HEMBB1002138	17.812	14.057	17.210	7.413	9. 287	10.613	20.319	9. 644
30	HEM381002139	51.267	37.549	168.617	27.467	17.855	27.091	16, 428	23. 177
50	HELS81002141	82.369	29.424	54. 387	14.566	15. 214	39.768	33. 139	22.856
	HEMBB1002142	70.553	42.309	156. 252	36.636	14. 797	26.769	15, 277	22.894
	HEMB81002145	40.661	16. 263	15.725	8. 229	13. 984	21.757	14.873	15. 525
	HEMB81002152	46. 728	36.893	105.608	65.422	40.064	25. 225	29. 211	42. 935
	HEM381002162	40. 153	34.008	96. 274	29.709	19.847	47.860	22.055	40. 550
35	HEMBB1002173	53. 191	41.151	147.055	26.912	34. 538	16.431	19.449	25. 327
33	HEM3B1002189	73.400	88.057	211.287	73.810	54. 029	46.682	45.749	55. 885
	HEMBB1002190	33. 242	51.561	233. 972	49. 809	19.665	27.376	13. 129	61.389
	HE19881002193	69.174	22.324	33.672	10.803	18. 423	27.938	24.748	16. 109
	HEMBB1002217	50. 175	37.602	98. 092	38.769	24. 723	33.043	18.735	39. 436
	HEMB81002218	596. 902	272.867	712.867	191.461	186. 314	373.711	195. 571	197.556
	HEMBB1002228	88. 583	45.763	205. 932	47.852	46. 693	41. 923	37.485	53.876
40	HEMB81002232	56.752	32.790	128.643	36.535	28. 693	32.710	31.447	41.940
	HEMBB1002245	31.084	9.332	17.943	11.049	11.834	11.864	17.012	14. 199
	HEMBB1002247	151.502	27.325	64. 167	10.018	26. 829	62.501	35.734	21.698
	HEL BB1002249	153. 327	94.814	380, 989	101.573	65. 579	80.049	62.653	85. 673
	HEY381002254	43.885	36.756	118.582	29.328	19. 323	11.675	12.693	22. 229
	HEY381002255	8.633	2.293	14. 174	8.771	1.813	2.385	3.358	3.589
45	HET 881002256	5.303	5.716	8, 530	6. 222	1.842	2.404	4.411	2. 295
	HE2881002271	160. 682	46.654	157.828	58. 291	63. 843	72.913	62.659	73.702
	HEY.881002280	24. 597	13.246	76.763	13.976	7.742	9.196	9. 200	16.479
	HE 881002296	67.004	21.270	52. 536	34. 388	49. 938	53.045	123.030	41.218
	REPBB1002300	94.815	28.682	50. 102	35. 939	13. 923	29.792	25. 246	21.629
	HE1981002302	51.059	31.157	28. 441	17. 568	17.905	26.026	22.516	30. 501
50	HELBB1002306			33,017				14. 490	
	HEY381002316	35.213	49.812		23.300	15.072	17. 296		16.293
		19.773	8.638	19.354	3.667	9.274	9.974	8.613	6.883
	HEL'8B1002326	201.896 85.792	126.797	406.052	154.628	89.356	85.970 29.959	54.052	98. 198
			48. 221	184. 126	47.724	32.764		17.415	34. 542
	HEMB81002329	69.191	21.714	43.746	25.618	17.775	24.892	32.481	27. 906
55	HEMBB1002340	18. 233	28.452	7.730	3.702	3. 055	4. 522	2.914	5.745

Table 48

	HEMB81002342	74.746	83. 579	169. 482	40.919	23. 495	26.453	33. 215	55. 420
	HEBB81002358	149.857	132. 962	286.214	85.160	50.855	67.646	36.624	78. 432
_	HEMB81002359	160.804	77.260	219.199	68.995	44.093	58.049	35. 955	51. 139
5	HEMBB1002364	102.885	74, 409	188.270	50.973	55.276	45.770	40.780	59.739
	HEMB81002366	152.074	77.016	248. 465	68.268	81.100	64.637	39. 912	60. 303
	HEMB81002371	44. 433	12. 342	26.565	13. 307	36.600	10. 553	9. 238	5. 351
	HEM881002381	134. 427	77. 953	207.310	57.210	48. 215	64.049	51.493	77.629
	HEMB81002383	164. 205	52, 312	94.064	31.346	31.368	30. 947	43.038	47.640
10	HEMBB1002387	196.859	164. 904	235.139	49, 485	25. 102	93.004	52.536	43. 092
	HEHBB1002409	82.986	49. 978	112.097	29. 207	15. 402	37.667	36.064	38. 132
	HEM881002413	123. 367	87. 690	361.106	87.505	57. 485	48.097 27.293	23. 254	49. 302 24. 444
	HEMB81002415	87.091	31.703	92.595	31.804	23.352	16. 980	18. 246	25, 253
	HEMBB1002424 HEMBB1002425	13. 162 84. 086	19.511 69.689	15. 995 238. 147	5.848 82.198	36.928	41.171	26.823	47. 957
	HEMBB1002427	143.727	26, 894	. 50. 430	25.865	40.707	52. 937	38.610	47. 517
15	HEMBB1002442	163.853	121. 153	501, 168	129.909	73. 231	81.033	47. 108	287. 238
	HEMBB1002447	107. 214	80.007	214.338	58.963	41.313	60.452	49. 159	44. 523
	HEHBB1002453	163. 250	93. 442	384.443	93.027	68.808	58. 565	45. 254	58.810
	HEMB81002457	116.756	104.520	330.657	83.026	46.720	50.971	38. 415	57.991
	HEMBB 1002458	18. 721	11.278	23. 232	9.587	7. 205	6.051	4.659	4. 343
	HEMB81002463	229.657	146.001	663.683	193.622	138.458	104. 827	52.827	110. 558
20	HEL'BB1002465	44. 210	23. 316	33.631	20.895	17. 932	26.471	19. 122	19.703
	HEM381002477	98. 948	27.813	153.875	11.062	36.071	16.072	13.791	8. 347
	HEP'881002479	23. 249	59.003	73.224	14.014	10.084	13.246	1.980 44.800	8. 949 75. 957
	HEMBB1002489 HEMBB1002492	78.748	24.690	71.038	31.400	39.869 1.453	5.606	1.415	2.381
	HEMBB1002495	9. 080 95. 752	6. 989 104. 949	26.130 301.328	60. 728	72.404	45. 161	24.771	61. 121
25	HEL:BB1002502	17. 132	17.866	14. 643	16. 170	15. 224	14.056	4.504	23.313
	HEM881002509	0.913	2. 235	7.269	4.304	0.743	1.283	1.504	6.154
	HEL'381002510	0.732	0.000	0.000	1.858	0.926	0.000	0.000	0.000
	HEMB81002520	249.875	127.604	585. 470	169. 423	138.712	90. 360	100. 598	112.828
	HEMBB1002522	24.741	27. 480	12.342	14.142	17. 452	5, 861	8. 292	8. 541
20	HEMBB1002527	63.012	61.066	87. 388 83. 329	46.392 21.750	29. 555 21. 479	37. 187 53. 227	25.642 440.333	36.089 38.710
30	HEMB81002530	72.655 40.398	45. 682 18. 832	10.308	9.953	5. 539	16. 743	11.880	8. 115
	HEMBB1002534	78. 552	49. 139	154.741	66.211	30.154	46.591	28.712	37, 112
	HEMB81002536	27.609	22. 843	52.264	17.646	8.234	13.078	23. 458	15.919
	HEMB81002544	24.012	6. 185	27.814	13.117	39. 363	15. 921	9.427	14.017
	HEM381002545	108. 234	31.929	243.949	50. 972	16.032	40. 343	31.828	13.472
35	HEYBB1002550	31.850	11.452	10.668	11.228	11.049	10, 100	14. 262	14. 910
	HEL'881002556	125. 621	89.607	311.607	79.974	50. 209	57.837 21.961	53.696 5.482	54.119 18.608
	HEMB81002571	33.047 75.252	21.526 55.132	54.457 229.479	14.847	25.892 31.521	43. 266	24.667	31.554
	HEMBB1002582	100. 572	56.574	258.453	63.093	45.740	39. 580	26.474	45. 912
	HEYBB1002584	8. 325	7.614	13.574	6.883	1.796	7.655	6.183	4. 955
40	HEE881002587	57.430	44. 383	60.900	47.981	30.048	30. 562	19, 161	20.854
,,,	HEE:881002590	114.241	78.587	179.926	65.737	28.629	43.657	33.101	34.032
	HEHBB1002596	278.617	90.944	275.018	69.006	68. 247	114. 505	88.149	59.750
	HEMB81002600	17.618	16.003	23.907	4. 699	9.726	10.133 37.423	7.945	8. 940 33. 698
	HE1381002601	67.910	48.188	183.948	45.346 36.733	38. 021 28. 849	35. 264	22.033	29.436
	HEMBB1002607	64. 941	36. 284	134.598	39. 424	22.220	31.501	15. 575	31.024
45	HEMB81002610	22.852	9. 200	51.294	16.832	6.664	12.856	6.433	6.515
	HEMB81002613	85. 026	60.872	161.891	47.532	36.559	44.841	24.569	31.062
	HEY 3B1002614	65.074	30.721	39.687	10.970	15. 910	13. 297	10.461	5. 438
	HEMB81002615	230.370	55. 581	35. 517	11.758	7.258	46.054	22.857	86.789
	HEMB81002617	69.016	67.288	254.296	42.530	30.217	36. 395	21.284	37.688
50	HEYB81002623	92.506	78124	204.116	60.739	20.110	48.078	32.253	43. 355
	HEMB1002624	77.755	27.026	163.976	33.209	25.309 7.219	20. 104 5. 864	6.990	24. 486 11. 537
	HEMBB1002631	10. 297 88. 049	18.892 68.172	12.879	41.853	40.290	23.649	21.781	44. 425
	HEL BB1002644	98. 956	65. 380	26.659	19. 268	9. 200	38.890	35.668	29.597
	HEP 881002654	127. 571	78.659	51.653	28.747	32.125	137.732	315.048	39.477
<i>==</i>	HEE'BB1002661	106. 501	46.651	47.116	19.470	20.684	30.561	24. 281	118.028
55									

Table 49

							- SER T	10 100
	100.783	42,600	100.008	36.841	24. 382	35.028	41.975	18.150
EE:881002664	179.828	131,008	395.057	51, 235	74.731	89, 130	40.814	55. 262
	2.206		5. 138	4. 981	2.314	4.033	3, 301	1.422
EMBB1002677		3.466						
EMBB1002683	118.247	69. 327	247.117	55. 886	44. 381	26.944	27.017	42.278
EMBB1002684	40.291	21, 056	46.317	17.772	9.039	5.460	8. 120	14.377
	30.893	12.882	26.031	19.059	3, 146	12.807	18.055	9, 131
EMBB1002686								
HEMB81002692	48.969	24. 335	52. 440	29.779	19.960	25.893	38.755	15.268
HEMB81002693	129.760	76.886	322,740	70.620	62, 314	67.760	73.429	39.005
ELBB1002697	41.673	38, 793	25. 105	8. 999	2.058	7,613	10.266	29.797
							79, 393	99. 532
HEMBB1002699	223.756	165.884		116.529	77.378	109.419		
HEMB81002702	13.506	15. 782	24. 367	3.561	6. 434	15.699	13. 253	24.914
HEMBB1002705	29. 934	20.276	16.478	21.230	7.599	11, 487	18, 202	30. 589
					13. 434	15.691	7.559	16.536
HEL:381002712	29.588	10.805	47.572	15.673				
IMR321000028	77.081	39. 937	40.934	18.725	8. 281	41.195	27. 733	21.472
IMR321000031	50.644	21.357	34.754	22.184	15.786	31.242	22.705	14. 148
IHR321000034	76.518	63. 230	37.290	51.243	23.808	43.858	26.605	67.455
								88, 100
1MR321000039	66.895	68.027	83.136	36.653	27, 339	62.232	57.760	
1MR321000044	1.614	0.000	0.000	0.000	0.000	0.000	1.970	0.000
IMR321000063	131.633	84. 822	66.499	84.753	43. 262	73.363	69, 831	80.878
				11.752	50.766	66.106	54. 160	47. 424
	157.704	34, 180	42.747					
MR321000089	52.645	22.980	31.408	17. 365	13.731	36.296	27. 222	10.181
IMR321000091	39.993	32,664	43.895	41.311	25.143	35.002	20, 444	63.906
		30, 112	69.445	15, 874	11.073	28. 505	106.044	24.660
LIVER1000004	45.674							11.282
LIVER1000008	23.703	14, 444	22.304	9.381	15.657	274.776	344. 333	
LIVER1000011	107.957	31, 187	106,032	30, 434	41.030	41.256	348.474	63.939
LIVER1000022	402.839	177.843	270.232	82.143	125. 292	206.780	141.934	124. 260
						33.045	34.820	42, 189
LIVER1000025	61.584	42.776	172.307	36.300	26.856			
LIVER1000030	62. 987	24.034	69.275	29.784	17.581	22. 393	51.178	22.556
LIVER1000045	27. 941	4.859	27, 468	7.384	9.755	14.426	20.651	24. 802
		117.998	24.240	23.527	16.373	7.456	27.795	66.724
LIVER1000046	180. 297			23.321				
LIVERIUUUU72	24.097	35.964	6.976	11.158	7.657	8.260	16.555	4. 898
LIVER1000077	90. 518	39, 165	17.306	13.193	25. 835	52.139	348.056	37.50
LIVER1000080	17.084	4, 918	5, 980	9, 600	2.294	5. 176	6, 495	4. 47
				18. 858	19. 278	176.018	481.085	27.74
LIVERIOOOOB6	82.711	55.169	150.708_					
LIVERTOOOGY	61.883	36.836	116.592	27.330	16.805	25. 266	35, 863	24. 16
LIVER1000095	54. 562	13,959	104, 146	23.878	13.158	200.163	137.395	5.50
LIVER1000097	138. 286	11.401	12.265	8, 127	9.389	9.669	32,751	7.15
							142.508	19.10
L1VER1000098	58. 055	39, 291	47.410	18. 991	19.124	20.338		
LIVER1000100	81.693	64, 546	94.504	29, 185	18.588	42.254	23.727	58.63
LIVER1000101	52. 507	16.303	57, 500	10. 286	8.662	17.642	6.129	27, 27
				11.568	9.377	13.216	102.126	16.90
LIVER1000106	46. 259	32. 121	32.438					
LIVER1000108	26. 277	50.565	62, 172	25. 422	16.619	17.243	38.369	18.50
LIVER1000115	23. 571	18.673	71.367	14. 244	11.023	17.910	427.626	11.13
LIVER1000120	100. 902	21.640	35. 183	16.565	26.236	39.037	87.151	16.24
					25.076	42.015	35, 937	23.83
LIVERICOO138	69. 624	27.584	56.479	22.794				
LIVER1000146	107.757	63.296	209.735	54, 534	42.231	45. 210	254, 168	42.46
LIVER1000148	141.467	42.327	108.510	37.031	31.920	62.584	125.466	65.72
	97. 282	37. 198	50.979	49. 952	35.021	43.954	52. 527	43, 22
11TO281A8A187 1								
LIVER1000157							00 100	30 7/
LIVER1000161	100.902	24.883	57.647	28. 329	31.562	42.781	89, 198	
			57.647 41.460				332.789	30.70
LIVER1000161	100. 902 97. 214	24.883 29.093	57.647 41.460	28. 329 25. 700	31.562 26.316	42.781 112.706	332.789	30.70
LIVER1000161 LIVER1000167 LIVER1000174	100. 902 97. 214 53. 927	24. 883 29. 093 23. 440	57.647 41.460 26.353	28. 329 25. 700 13. 595	31.562 26.316 12.625	42. 781 112. 706 36. 580	332.789 71.460	30.70 10.51
LIVER1000161 LIVER1000167 LIVER1000174 LIVER1000185	100. 902 97. 214 53. 927 49. 746	24.883 29.093 23.440 20.428	57.647 41.460 26.353 31.630	28. 329 25. 700 13. 595 13. 964	31. 562 26. 316 12. 625 13. 391	42.781 112.706 36.580 16.773	332.789 71.460 16.676	30.70 10.51 14.87
LIVER1000161 LIVER1000167 LIVER1000174 LIVER1000185 LIVER1000187	100. 902 97. 214 53. 927 49. 746 38. 332	24. 883 29. 093 23. 440 20. 428 8. 211	57.647 41.460 26.353 31.630 15.200	28. 329 25. 700 13. 595 13. 964 4. 654	31. 562 26. 316 12. 625 13. 391 8. 084	42.781 112.706 36.580 16.773 9.846	332.789 71.460 16.676 567.808	30.70 10.51 14.87 8.32
LIVER1000161 LIVER1000167 LIVER1000174 LIVER1000185	100. 902 97. 214 53. 927 49. 746 38. 332	24. 883 29. 093 23. 440 20. 428 8. 211	57.647 41.460 26.353 31.630	28. 329 25. 700 13. 595 13. 964	31. 562 26. 316 12. 625 13. 391	42.781 112.706 36.580 16.773	332.789 71.460 16.676 567.808 41.865	30.70 10.51 14.87 8.32
LIVER1000161 LIVER1000167 LIVER1000174 LIVER1000185 LIVER1000187 LIVER1000180	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672	24. 883 29. 093 23. 440 20. 428 8. 211 29. 635	57.647 41.460 26.353 31.630 15.200 50.518	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768	42.781 112.706 36.580 16.773 9.846 23.709	332.789 71.460 16.676 567.808 41.865	30.70 10.51 14.87 8.37
LIVERTOCO161 LIVERTOCO167 LIVERTOCO174 LIVERTOCO185 LIVERTOCO187 LIVERTOCO190 LIVERTOCO192	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875	24.883 29.093 23.440 20.428 8.211 29.635 53.337	57.647 41.460 26.353 31.630 15.200 50.518 99.330	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210	42.781 112.706 36.580 16.773 9.846 23.709 79.500	332.789 71.460 16.676 567.808 41.865 128.608	30.70 10.51 14.87 8.37 11.49 47.90
LIVERTOOD161 LIVERTOOD187 LIVERTOOD174 LIVERTOOD185 LIVERTOOD180 LIVERTOOD190 LIVERTOOD192 HAMMATOOD09	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036	24.883 29.093 23.440 20.428 8.211 29.635 53.337 77.266	57.647 41.460 26.353 31.630 15.200 50.518 99.330 234.005	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930	332.789 71.460 16.676 567.808 41.865 128.608 25.218	30. 70 10. 51 14. 87 8. 32 11. 49 47. 90 35. 90
LIVERTOCO161 LIVERTOCO167 LIVERTOCO174 LIVERTOCO185 LIVERTOCO187 LIVERTOCO190 LIVERTOCO192	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875	24.883 29.093 23.440 20.428 8.211 29.635 53.337	57.647 41.460 26.353 31.630 15.200 50.518 99.330	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014	30. 70 10. 51 14. 87 8. 37 11. 49 47. 90 35. 90 8. 11
LIVERTOOD161 LIVERTOOD167 LIVERTOOD174 LIVERTOOD185 LIVERTOOD185 LIVERTOOD189 LIVERTOOD190 LIVERTOOD190 HAPPATOOD18	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458	24.883 29.093 23.440 20.428 8.211 29.635 53.337 77.266 7.192	57.647 41.460 26.353 31.630 15.200 50.518 99.330 234.005 19.901	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930	332.789 71.460 16.676 567.808 41.865 128.608 25.218	30. 70 10. 51 14. 87 8. 32 11. 49 47. 90 35. 90 8. 11
LIVERTOOD161 LIVERTOOD167 LIVERTOOD174 LIVERTOOD185 LIVERTOOD185 LIVERTOOD189 LIVERTOOD192 HAPPATOOD189 HAPPATOOD189 HAPPATOOD189	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148	30. 70 10. 51 14. 87 8. 32 11. 49 47. 90 35. 90 8. 11
LIVERTOOD 61 LIVERTOOD 67 LIVERTOOD 74 LIVERTOOD 85 LIVERTOOD 87 LIVERTOOD 90 LIVERTOOD 90 LIVERTOOD 90 LIVERTOOD 90 LIVERTOOD 18	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999 58. 696	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927 30, 055	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049 181. 093	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037 40. 615	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18,169	30.70 10.5 14.8 8.32 11.49 47.90 35.90 8.11 30.1
LIVERTOOD161 LIVERTOOD167 LIVERTOOD174 LIVERTOOD185 LIVERTOOD185 LIVERTOOD189 LIVERTOOD192 HAPPATOOD189 HAPPATOOD189 HAPPATOOD189	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572 3. 468	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18.169 37.960	30.70 10.51 14.87 8.32 11.49 47.90 35.90 8.11 30.1 20.80 9.22
LIVERTOOD161 LIVERTOOD167 LIVERTOOD185 LIVERTOOD185 LIVERTOOD187 LIVERTOOD190 LIVERTOOD192 HAMMATOOD018 MAMMATOOD018 MAMMATOOD018 MAMMATOOD018 MAMMATOOD018 MAMMATOOD018	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999 58. 696 15. 610	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927 30, 055 5, 088	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049 181. 093 15. 411	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037 40. 615 7. 263	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572 3. 468	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18,169	30.70 10.51 14.86 8.32 11.49 47.90 35.90 8.11 30.1 20.80 9.2
LIVERTOOD161 LIVERTOOD167 LIVERTOOD185 LIVERTOOD185 LIVERTOOD187 LIVERTOOD189 LIVERTOOD190 LIVERTOOD192 HAMMATOOD018 MAMMATOOD018 MAMMATOOD018 MAMMATOOD024 MAMMATOOD024	100. 902 97. 214 53. 927 49. 745 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999 58. 696 15. 610 53. 706	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927 30, 055 5, 088 37, 358	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049 181. 093 15. 411 123. 944	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037 40. 615 7. 263 37. 766	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572 3. 468 29. 177	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176 11.662 24.650	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18.169 37.960 18.530	30. 70 10. 51 14. 87 8. 37 11. 49 47. 90 35. 90 8. 11 30. 1 20. 80 9. 2
LIVERTOOD161 LIVERTOOD174 LIVERTOOD174 LIVERTOOD185 LIVERTOOD185 LIVERTOOD185 LIVERTOOD185 LIVERTOOD195 LIVER	100. 902 97. 214 53. 927 49. 746 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999 58. 696 15. 610 53. 706 170. 220	24, 883 29, 093 23, 440 20, 428 8, 211 29, 635 53, 337 77, 266 7, 192 29, 927 30, 055 5, 088 37, 358 108, 774	57.647 41.460 26.353 31.630 50.518 99.330 234.005 19.901 150.049 181.093 15.411 123.944 290.077	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037 40. 615 7, 263 37. 766 126. 472	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572 3. 468 29. 177	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176 11.662 24.650 82.087	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18.169 37.960 18.530 70.843	30. 70 10. 51 14. 85 8. 32 11. 49 47. 90 8. 14 30. 1 20. 8 9. 2 21. 1 76. 2
LIVERTOOD161 LIVERTOOD167 LIVERTOOD185 LIVERTOOD185 LIVERTOOD187 LIVERTOOD189 LIVERTOOD192 HAMMATOOD018 MAMMATOOD18 MAMMATOOD18 MAMMATOOD18 MAMMATOOD18 MAMMATOOD18 MAMMATOOD18 MAMMATOOD24 MAMMATOOD25	100. 902 97. 214 53. 927 49. 745 38. 332 93. 672 141. 875 99. 036 40. 458 62. 999 58. 696 15. 610 53. 706	24.883 29.093 23.440 20.428 8.211 29.635 53.337 77.266 7.192 29.927 30.055 5.088 37.358	57. 647 41. 460 26. 353 31. 630 15. 200 50. 518 99. 330 234. 005 19. 901 150. 049 181. 093 15. 411 123. 944	28. 329 25. 700 13. 595 13. 964 4. 654 15. 812 32. 936 72. 924 13. 017 52. 037 40. 615 7. 263 37. 766	31. 562 26. 316 12. 625 13. 391 8. 084 18. 768 41. 210 40. 612 12. 921 36. 450 38. 572 3. 468 29. 177	42.781 112.706 36.580 16.773 9.846 23.709 79.500 44.930 18.315 42.958 34.176 11.662 24.650 82.087	332.789 71.460 16.676 567.808 41.865 128.608 25.218 13.014 38.148 18.169 37.960 18.530	30. 74 30. 70 10. 51 14. 87 8. 32 11. 43 47. 90 8. 18 30. 1 20. 86 9. 22 21. 1 76. 2 8. 8 8. 8

Table 50

						~	10.000	66 660
MALMA1000055	65.118	40.884	57.307	29.859	27.445	33. 405	22.066	23. 563
MALMA1000057	170.331	108.479	421, 160	100.366	84. 331	77. 475	42.047	55. 847
MALMA1000060	79.698	50.265	153, 319	49, 223	28, 927	42, 539	25.636	52. 458
MALMA 1 000069				48. 764	43.720	61. 342	45, 357	33.115
	118.921	35.010	182. 272					
HAPMA1000084	128. 354	92.819	277.404	87.542	63.176	65. 262	34.266	45.092
4824A1000085	40.199	20.019	40, 608	21.956	13, 181	18. 822	36.347	26. 209
HAPMA1000092	77.338	37. 915	167, 474	43.988	16, 101	26. 961	15.531	22.390
					11.893	44, 990	24.784	25. 160
MALYA1000096	55. 344	38. 495	38.888	25. 605				
MAPMA1000097	62.546	54.694	52. 522	52.269	24.807	65.730	25.787	23. 298
MAPMA1000102	67. 585	32.797	91,551	31.689	19.430	26.892	22.353	15.842
HALMA1000103	63.752	26. 301	89.530	30.004	12.188	31.709	11, 461	14.718
						24.686	16.649	17. 569
MAMMA 1000106	37. 915	23. 228	90.795	22.075	14. 445			
MAKMA1000117	58. 533	24. 502	43.190	22.445	16.140	27.418	15, 487	13.269
MAPMA1000118	104.168	58. 433	63.822	8,833	24. 039	42.731	38.062	43. 242
HALMA1000129	170.665	72,256	98.813	45, 970	22. 181	58.739	50.197	14, 587
HAPMA1000133		25.090	33.061	20.713	14. 310	34, 685	18,642	14. 101
	62.435							60.360
MAMMA1000134	106.522	79.090	245.344	90. 530	127. 758	76. 596	45, 325	
MANYA 1000139	78. 566	47.362	99.179	34. 535	22.772	37.601	28.841	28. 280
MAMMA1000141	30. 121	20. 5281	28. 150	13.910	5. 510	14.314	12, 120	15.748
MA#MA1000143	16.647	8.669	41.797	8.690	9. 949	10.059	4.040	8. 280
				28.777	86.623	42.827	51.840	42.986
MARMA1000150	128. 128	259.413	21.844					
MARMA 1000155	205. 031	88.642		110.884	80.817	97.755	63.045	78. 585
MAPMA 1000163	43. 543	36.898	57, 239	22.848	21.852	41.672	11.036	10.618
MAMMA 1000171	141.225	46. 928	265.746	98. 189	60.007	66.037	34.872	50.109
MANNA 1000173	103.027	21.955	68,080	33.572	25.668	45. 271	40, 340	52, 609
							5.974	4.015
MAMMA 1000175	19.316	8.683	7. 960	4. 550	3. 535	7.894		
MANSAA 1000183	57.490	35.830	148.702	42.892	23. 250	23.680	21.050	46.992
MALEMA 1000191	88.722	31.449	40.834	26.064	22. 392	26.766	36.253	27.729
MANSOA 1000 192	53.467	25.096	30, 205	28. 380	21.976	101. 288	128.339	44.025
MAMMA 1000 193	83, 936	36, 823	36.836	29.409	18.905	35, 131	35.059	36, 667
					49. 278	62. 924	38.858	66.720
MARMA 1000198	132. 127	93.550	347. 292	70.840				
HALMA 1000204	64. 455	59.079	71.789	26.771	29. 275	55. 156	62.132	49. 295
MANSAA 1000207	45.771	62.052	52.332	19.986	16.418	37.618	225.196	18.506
MANSKA 1000214	100.292	62.311	289. 223	62.541	32.825	57.748	32.755	39.770
HALMA 1000220	91.389	23.816	43.034	13.919	12.549	42.421	29, 143	20.494
					9. 426	18.802	27,741	17, 121
MAMMA 1000221	39. 338	35.655	11.931	39.315				
MANSAA 1000226	65.096	20. 174	11.901	11.838	17.236	23. 487	43.016	24.801
MAEMA 1000227	94. 333	64. 156	183.365	82.763	58.478	66.811	43. 961	53. 250
MANMA 1000230	116.378	47.908	97.869	47.218	38.196	56. 380	71.726	37.727
MANMA 1000241	53. 737	85.177	107.748	60.815	31.230	51.839	36.525	22.770
				144.478	51.682	86.017	93.183	198.398
MAKSMA 1000245	107.413	148.468	205. 437					
MAMMA 1000248	205. 478	88.411	342.827	76.468	51.702	110.723	70.650	60.978
MAMMA 1000251	115, 401	47.888	209.360	39.959	42.597	57.904	34. 572	51.015
HANSA 1000254	43.161	20.910	114.081	20.548	9.699	9. 885	5.346	32.024
MANSA 1000257	142.781	70.118	332.822	104. 425	84.387	124.673	78.270	116.10
				32.516	14.840	15. 513	23.805	35.51
LAMA 1000262	18. 952	34.301	19.786				22.616	37.03
MALAMA 1000264	59. 532	20.630	124.043	44.847	29.466	21.390		
MANMA 1000256	55. 476	28. 959	122.654	35.663	27.018	24. 021	20.212	38.28
MANSHA 1000270	142.968	64.234	270.948	75.022	64.760	68.130	64.006	73.99
MANMA 1000271	53.605	9,611	35, 682		16.139	24. 236	26.722	26.43
MAJSIA 1000277	56.407	16. 435	98. 448	19.751	12.725	33.047	23.839	33.01
		10.433					25, 204	
MAMMA 1 000 278	40. 286	13. 165	19.395	9.730	12.609	20. 423		22.23
MALMA 1000279	68.661	36.984	173.379	46.809	34.441	42.500	26. 143	48.59
MAKMA1000283	55. 199	27.095	46. 168	22.395	15.870	21.308	16. 298	18.50
MANMA 1000284	76.726	67.676	42.784	39.851	34.586	47.651	39, 169	48.34
MANIA 1000287		58.726	142.953	39, 301	31.007	27, 370	29,006	35.59
	73.583							
HALEA 1000294	457.450	361.106	313.407	116.696	112.848	343.951	155, 948	100.37
MAMMA1000298	31.731	25.511	41.413	16.220	16.320	14.676	22.043	20.20
	109.379	58.532	280.880	69. 156	44.790	36.788	28. 220	40.86
LIMEN TO LIBERTA	67. 505			11.073	33.859	26, 599	30, 177	30.81
MARMA 1000302		14. 147	18.804		12.695		13.353	15, 18
MARMA 1000303		1 10 44				14, 455	1 13.335	
MARMA 1000303 MARMA 1000305	32. 363	19.693	108.733	15. 375		44 144		1
MAMMA 1000303 MAMMA 1000305 MAMMA 1000307		75.098	397. 421	75.020	45. 244	68.757	131.117	
MAMMA 1000303 MAMMA 1000305 MAMMA 1000307	32.363 279.600	75.098	397. 421	75.020		68.757 8.895		
MARMA 1000303 MARMA 1000305	32. 363				45. 244		131.117	116.80 6.74 7.57

Table 51

ı	HAPMA1000313 [	70 644	60.660	64 317	10 741 1	CO (3C )	42 004	10 205	37. 303
	HAEMA1000313	79. 577 80. 910	69.550	54. 317	10.741	60.526	42.964	18. 206 18. 580	21. 385
	MAMMA 1000331		48.868	139.047	33.811	22.564	·15. 207		
		54.800	22.399	33.190	18. 244	16.273	30.688	26.611	30.790
	MAPSIA 1000339	69. 222	40.948	83.679	13.158	20.941	22.134	20.026	10.739
	MAMMA 1000340	57. 498	34. 708	164.968	32.922	28.610	23.069	18.858	23.519
	HALEMA 1000348	78.099	102.955	374.737	55.033	32. 546	66. 256	22. 303	23. 575
	HAE≌A1000356	152. 238	116.086	454.516	67.232	34. 525	47.884	22.865	61.267
	HAPMA1000358	34. 367	56.332	15. 362	15.091	15.743	17. 405	19.645	7. 358
İ	MAPMA 1000360	71.104	74.351	246.244	43.414	24.093	24. 945	14. 842	14. 739
	HAPMA 1000361	101.653	93.468	230.215	73. 577	45.022	37.236	37. 987	42. 992
	MAMMA 1000363	71.108	19.232	39.013	13.717	23.713	30.739	27.813	32. 485
	HAMMA1000370	171.867	108.830	110.466	80, 949	52.076	79.266	57, 877	247.810
	HAPMA1000371	100. 543	32.223	80.873	48.039	49.442	91.739	57.647	46. 599
	HAPMA 1000372	206.850	114. 326		130.138	79.980	80.890	54. 857	97. 509
	MARMA 1000385	72.074	60.911	238.462	40.061	34. 528	31.361	22. 458	45.681
	MALMA 1000388	118.855	69.094	105.789	42.626	50.059	55.389	37. 396	37.825
	MALMA1000395	97.031	44, 493	34, 493	20. 201	19,036	27.695	24. 269	17. 433
	MALMA 1000402	126.085	107. 637	256.584	68.415	45.669	61.486	30. 340	30.943
	MAKMA 1000403	87.558	63.749 (		64.857	45. 578	44.799	22.710	42. 239
	MANAMA 1000410	43.073	43. 539	94. 207	39.613	19.880	22.573	16. 272	21.003
	MANSIA 1000413	30.829	13.370	70.418	17.102	13.392 51.528	15. 291	11.599	15. 353
	MANSYA 1000414	125. 550	111.622	81.672	15.722	105.899	14. 549 121. 441	28. 214 55. 040	13.858 84.667
	MARMA1000416 MARMA1000421	179.864 131.712	103.793 73.475	427.214 307.780	107.383 70.841	55. 037	49, 498	34. 519	46. 482
	MAMMA 1000421	12,614	14. 628	307.780	16.100	11.675	22.441	18. 843	54, 831
	MANNA 1000422	34.100	22. 150	69.677	18, 461	13.815	15. 645	8. 500	8. 869
	MAMA1000424	9. 330	4. 056	35, 234	8. 171	0.971	2.769	0.745	7. 267
	MARIA1000429	575. 321	219. 603	317.414	158. 529	150.779	290. 300	196. 161	149.619
	MAISSA 1000423	143. 825	79. 993	275. 497	82.499	52.496	63. 425	43.337	66.733
	MARIA 1000432	65. 212	17. 117	24.472	28. 083	17.360	33. 881	27.547	29.615
	MANMA 1000437	89, 375	88. 947	265. 572	60.025	69.885	45. 195	30.823	31.510
	MANMA 1000444	120.017	124. 234		115.966	65. 200	66.888	31.943	88.274
	HAMMA1000446	50. 201	66.027	41.406	8.991	18.971	29.395	7. 985	37.220
	MALMA1000449	81.386	41.427	180.761	40.414	25. 983	35.232	23. 109	27.942
	MAIMA1000457	47.862	13.862	15.095	11.981	7.566	21.142	12.971	10.872
	MAMMA 1000458	34. 485	13.749	22.854	12.116	11, 199	18.881	15. 924	10.046
	MAMMA 1000468	8. 235	7.843	6.029	5.004	5.503	8.258	7. 138	1.618
	MAMA 1000472	250. 243	67.964	110.774	68.614	73.186	111.758	88.016	79.409
	MANSAA 1000473	54.174	16.506	40. 489	16.002	17.450	26.506	17. 741	13.900
	MAISSA 1000477	77.316	50. 237	238.943	56.460	38.807	32.776	36. 438	35. 332
	MAEMA 1000478	201.299	157.097		127.872	82.832	77, 444	49. 296	86.763
	MALMA1000483	107.340	74.564	252.463	60.824	31.055	44. 198	44. 157	87. 449
	MALSMA 1000490	14.473	14.068	16.023	12.496	8.202	15.654	11.091	12. 344
	MAEMA 1000498	32.756	10.554	20.693	10.676	19.830	19.282	13. 204	13.410
	HALEMA 1000500	23.016	17.584	49.151	15.706	13.914	19.063	11.094	22.904
	MANMA 1000501	196.637	102.490	468.793	104.118	67.761	83.834	76.446	86.912
	MAMMA 1000503	7.083	4.085	3.866	1.004	1.005	3.752	4.005	3. 248
	MARSA 1000506	201.452	116.279	151.434	56.847	78.502	149. 780	99. 352	54.069 33.127
	MARSA 1000510	70.898	18. 432	60.927	39. 187	33.327	42.829	40.993 19.339	17. 922
	MASSIA 1000515	43, 923	30, 031	85.637	35.744	18.805	34.061	19, 122	
	HAMA1000522	74.742 53.273	48.811 23.845	148.307	22.861	18.069	24.776	12.095	27.578
	MASSIA1000524	130.806	61.389	266. 529	71.558	50. 972	73.691	47. 484	55. 510
	MAMA1000528	38. 579	27.136	46, 940	35.839	15.860	29.316	19.300	24.797
	MASSMA1000534	32.603	20.088	33.950	10.973	7.185	10.580	7. 972	10.150
	MAPMA1000541	165, 518	58.806	85, 648	63.188	27.705	52.036	46. 200	39.018
	MAMMA1000550	119.597	203.059	41.184	24. 393	5.859	48. 433	766. 194	63.005
	MANMA 1000556	31.963	15.056	15. 588	8.634	11. 294	15. 698	21.467	16.597
•	MAMA1000559	57.738	31. 181	242, 155	29.443	19.030	26.908	13. 520	41.571
	MAMMA1000565	118.770	30.318	289.829	37.509	33.728	38.720	18.344	25.847
	MAMMA 1000567	77.050	44.379	224.645	48.804	41. 102	56.039	36.496	63.529
	MAYMA1000576	271.038	180.600	661.566	221.987	157.443	132.385	93.679	129.843
	MAMMA 1000582	54, 936	43.406	272.366	14. 342	18.896	29. 396	46.333	40.210
	HAMMA1000583	90.692	51.670	147.946	34.905	17. 175	23. 177	19.077	40.824

Table 52

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	HAMMA1000585	89.865	50.008	288.673	52. 259	29. 243	39.188	24.088	45.734
			14.789	58. 279	12.415	6. 584	14.410	15.734	6.826
	HAEMA1000587	47.955			28.019	20.094	28.578	24. 299	19. 949
5	MAMMA1000591	77.705	38. 280	81.784		59. 244	55. 681	43.577	75. 029
	MAEMA 1000594	194. 593	94. 384	488. 898	91.064			199. 968	160. 426
	MALYA1000597	496. 923	264. 906		196. 294	121. 483	306.397	97.598	149. 183
	MAMMA 1000605	324. 584	183.667		209. 555	135. 844	158.096		
	MAPMA1000612	68.113	22.051	42.999	14.074	19. 294	41.220	29.460	15.713
	MAYMA1000614	580.099	136.874	402.890	69.022	127.808	309.892	249.344	
10	MAPMA1000616	2. 590	16.442	13.809	1.109	3. 011	7.500	3.036	3.188
. •	MAMMA1000621	19. 258	12.723	14.307	13. 200	5. 971	12.028	11.561	11.081
	MAMMA 1000623	60.189	23. 285	25.913	12.057	10.648	23. 327	19. 218	20.667
	MALMA1000625	651.334	249.117	346.876	155. 944	192.671	373.924	300.473	274. 263
	MAKMA1000635	4. 459	2. 994	4.756	2.883	0.000	4. 118	5. 584	9. 542
	MAK-4A1000643	24, 259	51.698	115, 511	47.881	17.554	52.330	16.308	38. 448
15	HAPMA1000546	72.487	111, 121	22.868	9. 213	27.074	81.604	46.859	34.048
13	MALYA1000652	152. 920	94, 568	319.943	76.610	67.817	87.605	41.747	77.720
	MARMA1000657	116.830	41.097	278.504	38. 131	36. 289	67.327	34. 224	32.593
	HAMMA 1000664	48. 908	37.993	133.863	26.712	16.308	21, 135	14. 102	35. 215
	MAKMA 1000667	77. 285	24. 312	99.732	25.027	29. 493	43.769	22. 193	24. 502
	MAMMA1000668	42. 561	28.100	-54.970	17.454	18. 336	50.398	38.233	26.553
•	MALMA1000669	22.797	14. 382	57.803	14.670	6. 337	12.841	7.392	12.088
20	MAKMA1000670	66.748	22.566	46.836	26.498	25.826	33. 332	38.768	39. 130
	MACMA1000672	128. 331	25. 209	67.913	35. 262	28. 783	64.713	38.934	40. 592
	MAMMA1000681	66. 397	40.677	32.249	14. 404	13. 181	26.710	30.054	37.369
	MARSMA1000684	85.908	107. 381	66.100	35. 992	32.881	41.006	36.719	77.834
	MAMMA 1 0006 9 6	165. 293	107. 442	551.458	130.714	88.510	70.985	43.857	55. 551
	MAMMA1000702	82. 316	25. 689	52.797	22.639	22.884	48.899	39.297	29. 636
25	MARMA1000706	81.416	25. 442	34.529	20. 432	15.562	39.909	33.303	25. 371
	MARSA 1000707	128. 277	17. 100	51.835	15.001	33.473	48.628	46.555	24.075
	MAMMA1000713	75. 263	59.677	109.995	37. 970	23. 975	33.874	30.149	39. 491
	MARSAA1000714	228. 366	288.017	246.261	56.045	25. 380	80.480	51.219	64. 589 49. 032
	MAMA1000718	98, 208	92.149	245.750	79.940	49.064	50.180	40. 223 29. 904	60.252
	MANSA 1000720	158.737	111.227	446.586	101.175	73.612	78.021	18.845	24.784
30	MAMMA1000723	64. 930	49.053	148.286	40.276	28. 806	19.434	7.536	7.367
	MANMA1000731	31.516	11.357	68.834	12.436	11.755	7. 989	35.440	49. 335
	MAMMA1000732	121. 291	56.513	230.064	68.746	51.582	14.100	8.586	10.632
	MAMMA1000733	24. 525	14.171	58.717	16.852	7.153	84. 456	43.098	98.011
	MARMA1000734	113.011	127.466	142.152	102.345	44.860	73.418	82.810	69. 461
	MAMA 1000735	142.978	48.490	130.520	34. 595 38. 642	40.252 18.942	31.735	48. 926	35. 128
35	MAMA1000738	110.304	61.504	28.831 281.287	76. 261	79.000	63.977	43. 557	40. 380
	PANMA1000744	140. 264 26. 385	94.669 50.110	37. 264	16. 895	10.790	35. 280	3. 177	11.010
	MAMMA1000746		36.619	52.587	30. 957	36.810	46.899	25.359	24.846
	CASSIA 1000748	73.879 42.505	27.882	58. 087	44. 924	28.537	43.075	32.581	61.052
	MANMA1000752	55.785	55.799	193. 100	53.436	25.798	29.655	21.969	44. 384
	MAIMA1000757	314.709	210.647	536. 246	187.416	161.327	151.926	112.625	152.076
40	MARMA1000760	218. 937	178.377	534. 346	131.736	100.173	95. 443	58. 158	91.220
	MAMA1000761	147. 993	73.793	349.399	85. 319	65.436	75.180	43.310	63.428
	MAMA1000775	75.873	25.684	170.040	34. 150	30.063	20.938	15.825	18. 992
	MANA1000776	101.205	81.986	253. 211	57.436	51.043	51.597	28. 394	33.452
	MAINA1000778	71.839	47.596	214.100	42,749	28.124	29.701	17.866	26.497
	MALMA1000781	67.901	30.437	97.580	26.658	23.265	29.056	17.488	26.972
45	MANA1000782	285.062	65.796	174.951	84.753	88.062	151.891	90.446	86.369
40	MANMA1000784	135.655	91.366	264.154	67.248	65. 127	26.625	29.991	78.501
	MANSA1000788	143.478	49.979	98. 983	34. 503	30.600	55.026	29.032	46.210
	MAMMA1000798	62.822	41.315	139.860		26.873	27.100	11.942	32.539
	MAMMA1000802	132.633	86.328	341.638		64, 234	64.772	38. 532	61.561
	MANMA 1000810	150.779	88. 200	372.241	99. 538	80.592	81.887	42.150	57.891
50	MAMMA1000813	31.571	14.636	31.497		9.356		12.633	10.718
50	MACCA1000814	197.602	134. 253	279. 885				64.626	62.091
	MALMA1000824	65, 693	21.602	64.020		35.405		31.671	38.813
	MANSA 1000827	146, 098	70.894	157.448				33.051	44.519
	MALMA1000831	55, 332	19.954	29.847				16.602	6.497
	MALMA1000838	39. 583	28.962	39.815		49. 251	39.669		19. 273
	Fran 1000030	1 99. 303	1 20.302		1 23. 001				

Table 53

	1440141000000 T	167 000 1	110 400 1	700 OCA	111 004 1	06 640 1	05 054 I	57, 386	102.963
	MAKYA1000839	157.898	138.792		113.084	86.648	85. 964		
	MAKMA1000841	44.843	37.288	50.074	28.351	19.319	37. 537	13.012	20. 655
5	MAKMA 1000842	174. 347	36.747	169.008	44.926	48.610	78. 492	50.804	35. 389
3	MALMA1000843	8. 643	4.650	14.084	4.758	2. 185	6. 547	5. 283	1.757
	MACHA1000845	40,044	33, 955	33.012	21.488	15.747	23.310	17.728	15.607
	HAMMA1000851	197.033	79.321	307.054	96.446	73.025	75, 853	98. 526	72.039
	HAL-1000854	66.648	33. 221	63.298	17.429	20. 157	33. 288	22, 320	21.685
	MALMA1000855	10, 264	4. 185	17.702	3.794	3. 995	2. 454	9, 158	3. 568
						38.378	82.629	60. 529	25.726
10	MAMMA1000856	186. 269	40.945	84.561	27. 973			20. 320	33.032
	MALMA1000859	64. 234	121.939	60.662	34.958	42.945	39. 557		
	MALMA1000862	40. 107	21, 345	23.693	16.808	28. 277	22. 561	14. 238	14.693
	MAPMA1000863	98. 576	70.285	234. 996	67.796	55. 216	72.466	36.802	70.885
	HAMMA1000865	1.106	0.000	0.000	0.000	2.321	0.000	0.000	0.000
	MAPMA1000867	46. 228	24.216	64.376	21.736	17.699	18.758	10.742	6.954
	MAPYA1000875	124.814	80.537	231, 558	88.627	57.015	82.859	46.826	53.611
15	MALYA1000876	87,475	36.523	94. 191	19.763	21.466	42.434	27, 201	24. 439
	MALMA 1000877	201.968	107.716	538. 232	164.333	86.827	114. 380	80.171	97.872
				257.022	71.323	29.066	47. 487	36.714	37.365
	MALMA1000878	99.671	67.833				44. 996	19. 238	35. 353
	HAMMA1000880	76.396	60.884	153. 335	45.836	17.649			38.045
	MARCHA1000881	63.646	33.072	177. 731	43.034	30.410	31.086	12. 184	
	MALMA1000883	71.807	24. 931	43, 109	16.630	18.675	40. 320	44. 419	55. 440
20	MAMMA1000897	88.466	0.000	7.404	0.000	0.000	0.000	0.000	0.721
	MAMMA1000898	380.818	62.977	134.846	45, 311	63, 221	164. 332	122,071	52.933
	MAIMA1000905	97.555	63. 528	161.117	57.777	42. 205	50.312	28. 216	42.710
	MAMMA 1 000906	57.788	33.146	125.096	29.019	13.531	29. 380	16.982	14. 930
	MANMA1000908	30, 597	19. 222	40.351	11.584	5. 445	10. 392	13.469	11.612
	MAMMA1000911	9. 952	29. 425	3. 998	9. 963	1.886	7.419	5. 350	126.406
25	MAYMA1000914	82.184	23. 137	69. 228	20.659	18.111	35. 329	22.616	18.859
25	MANMA1000920	92. 123	62.032	37, 206	16.675	15. 550	47. 235	47.680	26.801
	MALMA1000921	107.169	69, 026	207.821	102. 347	60, 403	64.787	35. 902	77.424
	MAMA1000931	211, 796	140, 234	424.498	95.390	40. 229	51.643	49. 349	95.211
	HAMMA1000940	145. 411	82. 982	268.876	70.972	55. 532	61.420	51.119	60.328
				509.857	131, 193	79.478	106.717	53. 292	91.187
	MAIMA1000941	182.800	134. 847			68. 234	90.801	63. 506	75.814
30	MARMA1000942	195.078	123, 131	446. 428	117.435			51.063	85. 539
	MACMA1000943	196. 926	99.988	558.754	109.551	89.006	81.092		
	MANSA 1000952	161.019	97.081	355. 265	78.330	98.779	104. 172	79.021	96.980
	MAINA1000956	43. 741	16.217	14. 918	11.103	5. 840	41. 230	24. 471	6.893
	MAZELA 1000957	95. 532	53.066	225.645	64.794	42.610	47. 323	34. 337	45. 567
	MANMA 1 000962	281.600	192.048	781.968	204.962	120.611	123. 900	84.354	140.995
	MANMA1000966	151.087	157.558	417.591	111.282	64.746	81.685	51.694	78.953
35	MANMA 1 000968	217.975	107.043	313.251	58. 469	41.964	45.044	41.392	63.998
	MANMA1000972	18. 150	48.148	119, 482	22.427	18.041	15.672	12.870	13. 135
	MANMA1000973	36.667	18.879	24.787	11.758	12.527	19.441	17.828	22.312
	MANSIA 1000975	44. 972	19.058	38.995	20.137	30.793	22.864	65.817	45.398
	MAMMA 1000976	122.625	67.075	216.981	70.671	50, 470	91.475	60.614	81.173
	HAMMA1000979	81.812	102.452	145, 415	68. 435	53.443	56. 902	38.749	89.759
40	HAMMA1000986	118. 211	39.368	239.204	68.513	49.208	56.431	42.354	94. 152
, ,	HAMMA1000987	81.466	50.679	249.660	43.686	35, 580	49.753	23.004	41.997
	MAL®MA1000988	150. 907	68, 191	242.562	63.946	34. 252	81. 162	48. 528	86.723
					21, 154	26. 136	49. 152	44. 373	50. 523
	MANMA 1000994	101. 984	21.000	41.248		105.673	107. 213	56.957	84. 216
	MAMMA1000998	166.669	75. 193	367.111	91.202		35. 674	26. 101	59.032
	MANMA 1001003	73.580	37.252	146, 092	47.279	34. 315			0.000
45	MAPEN 1001007	3.055	0.000	5. 547	0.000	1.411	3. 633	0.800	
	MAYMA1001008	40. 892	31.048	65. 220	38.501	74.831	38. 859	47.979	31. 121
	MANSMA 1001013	135. 486	126.855	372.544	93.280	57.270	56.674	44.237	52.328
	MAMMA 1001014	85. 681	25. 361	77.414	32.516	25. 227	20. 809	35. 346	16.624
	MANMA 1001021	93.867	49. 224	180.659	41.205	34.542	34. 975	35. 352	29.726
	MAMMA 1001024	141.736	49.918	229. 735	52.670	41.069	54. 541	41.726	36.711
50	MANMA 1001025	13.661	8.964	12.310	5.843	13.733	6.698	4. 305	5.091
50	MAPTA 1001028	36. 353	24.719	14.061	10.363	34.518	16. 233	15.746	11.316
	HAPMA1001030	33. 596	27.602	35. 295	20. 296	15.861	14. 989	25.031	23.535
	MAMMA 1001035	235. 880	125.555	517.898	181.208	139, 149	129.655	96. 375	134.509
	HAPMA1001036		45.689	152.344		47, 114	60. 433	40.803	40.973
	ALIEUTIA WILLIAM	1 133. 350	1 43.003	136.344	100.032				
	HAPYA 1001037	180.875	100.457	403, 651	52.277	55, 761	72. 026	38.313	51.826

Table 54

41	MAPMA 1001038	26.248	12.160	150.692	32.729	5. 309	10.436	15. 903	27.263
	MAL'A1001041	113. 237	27.602	43.846	32.708	45. 924	47.820	46.929	16,614
	MALYA 1001043	218.483	23.847	68. 163	22. 306	10.449	41.046	45.779	31.087
	MAPMA 1001050	157.361	80.096		71. 548	69.197	49.684	13. 493	49, 872
					63. 324	43. 343	21. 184	38. 007	39.478
	MAMMA 1001054	102.456	62.728	134.003	52. 931	26.061	111. 283	69.714	40.010
	MAMMA 1001059	136.357	48. 942	59. 998	140. 850	119.334	176. 295	158. 563	60. 324
	MAMMA 1001066	387.798						19, 782	
	MAMMA 1001067	82.327	39. 420	127.017	37.076	29.891	30.670		14. 257
)	MAPPA 1001072	150.398	31.601	52.273	21. 983	32.143	57. 421	47. 051	26. 375
	MAMMA 1001073	101.957	23. 218	17.217	11. 405	43. 228	24.053	24. 142	5. 176
	MAPS 4 1001074	104. 201	41.827	240. 332	94. 124	56.071	89.717	16. 387	14.966
	MAZYA 100 1075	32.081	34.601	23.705	29. 782	21, 196	23. 184	14.757	17.497
	MARMA1001078	102.185	111. 402	317.478	75.869	35.841	49.660	67. 285	67. 244
	MALMA1001080	367.248	210.764	130.259	89.003	81.982	185.406	141.739	266. 507
	MAPYA 1001082	50.264	39.773	20.039	17.602	43.163	26. 358	17. 452	14. 352
'	MACCAMINATION 1091	3.576	11.403	27. 522	0.000	18. 321	4. 593	0.000	0.000
	MAMMA 1001092	50.554	25.306	48. 577	16. 425	15. 153	18.849	11.524	4. 155
	MACMA 1001094	353. 180	72. 506	112.379	42. 145	78.386	130.368	113.824	62.964
	MAMMA 1001 105	138.777	111. 2261	113. 121	82.426	80.960	45. 158	16.891	45.652
	MAMMA 1001110	15.141	8.661	7.407	3. 823	5. 537	6.280	3. 216	4. 392
	MAMMA 1001126	299.120	223.060		194. 522	164.920	119.375	96. 413	88.784
)	MAMMA1001133	243.826	187.024		144. 907	119.301	111.573	67. 515	94. 605
	MAMMA 1001 139	291.212	867.784	447.960	473. 187	227.579	348.627	121. 382	173.640
	MAMA1001141	36.320	18. 295	40.066	9.930	5. 202	26.277	16. 337	13. 996
	MANNA 1001143	163.308	70. 387	153.588	67.249	59.919	67.023	43.805	40. 903
	MANMA1001145	110.718	43.148	141.067	30. 89Q	31.851	11.000	10, 119	13. 322
	MANSUA 100 1150	80.076	29.005	50. 289	15. 249	7.495	33.674	48. 052	22.629
5	MACMA1001154	203.206	129.777	429.878	121.700	90.014	77.333	45. 155	71.154
	MANMA1001159	46.847	28.763	19.301	13.704	8. 444	23.404	21.864	24. 248
	MARMA1001161	185.601	233.229	485. 605	141. 151	109.607	107. 154	96. 161	79.043
	MATERIA 1001162	196. 299	51.198	67.587	29.962	40.684	78.949	43. 247	18.714
	MAPAMA 1001181	116.505	35.688	88.127	33.728	40.701	41. 280	16.749	26.312
	MAMMA 1001185	155.118	85. 120	303.506	69.532	51.017	85. 296	42.211	48. 082
,	MAMMA 1001189	60.587	31.052	16.618	30. 386	22.337	29.809	50.065	54.044
	MANA 1001191	120, 521	18.093	41.909	22.249	21.661	39. 122	50.157	24.623
	MAMA1001198	229.338	561.556	755. 924	695.028	205.811	536.623	412.766	746.035
	MANMA1001202	322, 950	274.854	664.569	248.672	218.550	168. 136	144. 829	179.567
	MAMMATOO1203	170, 551	101.121	330.599	85. 243	72. 915	53. 390	44. 564	52. 183
	MANNA1001205	132.103	114.504	202.256	65. 195	71.217	61.327	43.601	48. 988
	MAMMA1001208	55.417	28. 101	30.608	21.282	25.686	27. 394	20.015	15. 433
	MAMMA1001215	199.721	123.016	194.852	82.919	72.839	87.841	68. 245	60.078
	MAMMA 1001220	223. 133	154.557	404.346	110.968	91.387	74.073	58.534	62.841
	MARMA1001222	5. 585	4. 936	6.763	1.952	0.474	2. 171	20.800	5.022
	MANMA1001223	94.809	29.294	42.345	15.601	20.861	20.316	32.445	15.726
	CAMMA1001232	130, 199	45.692	227. 125	47.671	38.837	45, 692	59.906	32.862
	MARMA 1001234	129.344	27.935	227.692	95.815	64.344	61.799	49.210	34.673
1	MANAMA1001237	29.560	11.083	23.224	7. 241	4.489	20.199	16.883	11.003
	MANUA 1001243	20.832	11.598	47.127	7.253	32.689	20.073	7.954	6.544
	MANNA 1001244	44. 925	10.751	11.473	9.770	11.102	14.902	16.779	4. 470
	MANMA1001249	43.758	23.671	15.616	19.023	10.556	26.846	10.975	13.758
	MAMMA 1001256	169.303	81.917		187.649	131.656	44.850	55.325	59.786
	MANMA1001259	70.213	24.036	18.445	18.447	25. 202	45. 289		
i	MAMMA1001260	154. 426	64.153	81.115	52.438	46.566	80.874	64. 937	87.761
	MANMA1001252	153.326	53.618	54.054	40.354	54. 252	66.416	134. 449	25.835
	MANEMA 1001268	97.760	53.599	146.494	47.068	42.826	34.360	20. 976	28. 286
	MAL=4A1001271	305.116	66.364	106.518	32.761	65.392	128.314	130.796	39. 913
	HALEMA1001274	73, 329	94. 857	235. 488	85.814	64. 385	71.860	51.097	62.114
	MAMMA1001280	66. 399	17.595	13.218	9.853	3.831	37.015	12.303	6.374
1	MANNA 1001283	145. 535	67.060	129. 301	56.055	38. 490	56. 197	52.661	34.076
	MALMA 1001284	253.434	60.199	204. 903	48.719	63.272	100.485	93.658	76.590
	MANIMA 1001286	86.284	38.290	49. 421	32. 175	40.490	57.666	59.470	32.210
	MANMA 1001289	169.737	90.053	62.200	32. 142	102.670	66.398	64.913	47.082
	MAP 441001292	103.898	20.400	28.796	15. 498	31.006	29.378	26.545	31.970
	HAMMA1001296	225.022	173.717	324, 251	133.662	60.125	88. 173	70.926	89.316
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Table 55

	MAUNIATA T	00 070	60 100	040 000	30 (05 ]	12 020	26 676 1	27 022 1	27.836
	MAK-4A1001298	80.876	60.189	230.669	38. 485	32.838	36.675	27.032	
	HAMMA1001305	153. 258	67.563	147.529	36. 286	31.766	65.281	41.627	30. 730
5	HAMMA1001309	6.490	8.306	6. 534	3.627	4.269	0.000	5, 861	6. 705
· ·	MAPMA 1001310	148. 253	53.093	165. 786	46.753	41. 171	63.488	82.639	54. 927
	MALMA1001322	20.005	14.809	29, 403	19.332	11.227	14. 549	14. 163	15.700
	MAMMA1001324	82.605	28.652	85. 996	52.506	31.339	47.688	30.365	20.779
	MAMMA1001330	180. 949	117,040	245. 119	52, 680	15. 121	97.891	81, 121	27.980
	MAMMA1001333	101.707	75, 972	213. 812	59.950	49.965	59.640	32.340	37. 307
	MAMMA1001334	156.564	108.340	81.315	54.901	34.949	73.570	65. 555	73. 287
10		105. 507				20. 426	44. 148	21.930	33.068
	MAUMA1001337		35.111	33.563	17. 119			42. 286	29. 671
	MAPMA1001341	100.751	32.100	79.257	23.788	38.019	38.614		
	MAPMA1001343	128.875	95. 425	301.822	74.316	77.337	85. 437	18. 963	98. 899
	MALTHA1001344	32.880	35. 930	40.648	21.963	23.320	30.315	15.394	27. 074
	MAPMA1001346	49.749	17. 537	51.635	21.147	20.480	22. 107	26. 805	24. 306
.0	MALMA1001383	202.565	186.453	597.532	117.676	100.238	103.083	68.993	76.274
15	88E1001388	149. 105	66.100	213.624	45.488	52.686	66.868	85. 346	57. 974
	MALMA1001396	197. 435	81.919	430. 433	80.848	94.812	95, 399	75. 293	90.889
	MARMA1001397	116. 167	86.809	175, 125	67.323	58.676	56.833	61.558	52. 233
	MALMA 1001401	101.761	72.090;	194.999	62.960	48. 162	57.422	73, 403	78.023
	MANYA1001408	62.875	17.757	62.603	9.779	13.557	44. 301	11.008	20. 408
	MALMA1001411	271.344	54. 507	67. 489	20.558	68.557	157.085	134. 884	38. 338
20	MAPMA1001414				27.219	20.603	32.791	16.798	25. 126
20		74. 836	21.511	88. 459			89. 184	99. 527	41.848
•	MAMMA1001415	207.635	38. 228	51.690	26.716	68.700		31. 339	23, 195
	MAMA1001418	103.090	36. 102	91.976	39. 234	28. 949	27.016		
	MACMA1001419	106.299	52. 357	210.943	52.570	45. 256	41.351	37.624	25. 914
	MAMMA1001420	133. 835	25. 587	149, 981	15.816	19.703	28.670	26. 323	15.896
	MALMA1001426	265. 539	180.062	165.308	87.320	89.096	170.869	109.848	84.772
25	MANUA 1001428	310.313	180. 134	229.960	136.337	147.398	262. 499	135. 345	83.047
	MAZMA1001432	266. 375	107.317	387.676	85.786	60. 159	83.974	37. 205	60.775
	MALMA1001435	99. 596	48.079	193.151	53.623	27. 154	41.869	30. 388	39.835
	MAPPA 1001442	103.071	100.872	193. 544	78.030	54.054	54. 359	43.164	50.728
	MALMA 1001446	180.367	105, 551	197.748	98. 484	72.694	46.485	39.641	61.589
	MANSAA 100 1450	67.785	51.961	68.660	34, 362	32. 591	32.211	28. 904	9. 424
00	MAMMA 1001452	180.732	124. 244	432.438	115.549	111.829	104. 153	92.517	96.081
30									
	IMANSMATANTIAS S	I 57R 58R	255 549	770 820	1359 206	1 364, 762	388, 404	1 209, 219	264.053
	MARSIA 1001465	528. 588	255. 549	770.820	359. 206	364.762 17 988	388. 404 24. 584	209.219	264. 053 17. 981
	MAMMA1001476	33.639	19.551	25. 289	5. 909	17.988	24.584	26, 252	17. 981
	MAMMA1001476	33.639 117.183	19.551 61.333	25. 289 147. 393	5. 909 46. 785	17. 988 39. 649	24.584 32.143	26. 252 33. 776	17. 981 40. 723
	MASSA 1001476 MASSA 1001478 HASSA 1001479	33.639 117.183 156.131	19.551 61.333 59.931	25. 289 147. 393 31. 646	5. 909 46. 785 28. 808	17. 988 39. 649 44. 671	24.584 32.143 62.901	26. 252 33. 776 69. 911	17. 981 40. 723 26. 759
	MAMMA1001476 MAMMA1001478 HAMMA1001479 MAMMA1001487	33.639 117.183 156.131 67.613	19.551 61.333 59.931 53.042	25. 289 147. 393 31. 646 92. 480	5. 909 46. 785 28. 808 34. 978	17. 988 39. 649 44. 671 30. 928	24.584 32.143 62.901 40.427	26, 252 33, 776 69, 911 27, 489	17. 981 40. 723 26. 759 11. 238
35	MAMMA1001476 MAMMA1001478 MAMMA1001479 MAMMA1001487 MAMMA1001498	33.639 117.183 156.131 67.613 96.522	19.551 61.333 59.931 53.042 111.213	25. 289 147. 393 31. 646 92. 480 222. 159	5. 909 46. 785 28. 808 34. 978 50. 813	17. 988 39. 649 44. 671 30. 928 14. 811	24.584 32.143 62.901 40.427 23.385	26. 252 33. 776 69. 911 27. 489 56. 209	17. 981 40. 723 26. 759 11. 238 28. 054
35	MAMA 1001476 MAMA 1001478 MAMA 1001479 MAMA 1001487 MAMA 1001488 MAMA 1001501	33.639 117.183 156.131 67.613 96.522 216.969	19.551 61.333 59.931 53.042 111.213 55.879	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459	5. 909 46. 785 28. 808 34. 978 50. 813 38. 369	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036
35	MAMA 1001 476 MAMA 1001 478 HAMMA 1001 479 HAMMA 1001 479 HAMMA 1001 498 HAMMA 1001 501 MAMA 1001 501	33.639 117.183 156.131 67.613 96.522 216.969 124.674	19.551 61.333 59.931 51.042 111.213 55.879 57.815	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281	5. 909 46. 785 28. 808 34. 978 50. 813 38. 369 46. 452	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860
35	MAMATOD1476 MAMATOD1478 MAMATOD1479 MAMATOD1479 MAMATOD1498 MAMATOD1501 MAMATOD1501 MAMATOD1501 MAMATOD1502 MAMATOD1510	33.639 117.183 156.131 67.613 96.522 216.969 124.674 27.993	19.551 61.333 59.931 53.042 111.213 55.879 57.815 7.591	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577	5. 909 46. 785 28. 808 34. 978 50. 813 38. 369 46. 452 10. 197	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048
35	MAMMATOD1476 MAMMATOD1478 MAMMATOD1479 MAMMATOD1409 MAMMATOD1501 MAMMATOD1501 MAMMATOD1501 MAMMATOD1502 MAMMATOD1510 MAMMATOD1522	33, 639 117, 183 156, 131 67, 613 96, 522 216, 969 124, 674 27, 993 56, 601	19.551 61.333 59.931 53.042 111.213 55.879 57.815 7.591 24.819	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994	26, 252 33, 776 69, 911 27, 489 56, 209 43, 395 34, 762 14, 922 29, 481	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416
35	MAMMATOD1476 MAMMATOD1478 MAMMATOD1479 MAMMATOD1499 MAMMATOD1499 MAMMATOD1502 MAMMATOD1510 MAMMATOD1510 MAMMATOD1522 MAMMATOD1529	33, 639 117, 183 156, 131 67, 613 96, 522 216, 969 124, 674 27, 993 56, 601 83, 190	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170	26, 252 33, 776 69, 911 27, 489 56, 209 43, 395 34, 762 14, 922 29, 481 29, 923	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596
	MAMMATOD1476 MAMMATOD1478 MAMMATOD1479 MAMMATOD1409 MAMMATOD1501 MAMMATOD1501 MAMMATOD1501 MAMMATOD1502 MAMMATOD1510 MAMMATOD1522	33, 639 117, 183 156, 131 67, 613 96, 522 216, 969 124, 674 27, 993 56, 601	19.551 61.333 59.931 53.042 111.213 55.879 57.815 7.591 24.819	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522	26, 252 33, 776 69, 911 27, 489 56, 209 43, 395 34, 762 14, 922 29, 481 29, 923 25, 583	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896
35	MAMMATOD1476 MAMMATOD1478 MAMMATOD1479 MAMMATOD1499 MAMMATOD1499 MAMMATOD1502 MAMMATOD1510 MAMMATOD1510 MAMMATOD1522 MAMMATOD1529	33, 639 117, 183 156, 131 67, 613 96, 522 216, 969 124, 674 27, 993 56, 601 83, 190	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298	26, 252 33, 776 69, 911 27, 489 56, 209 43, 395 34, 762 14, 922 29, 481 29, 923 25, 583 46, 390	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233
	MAMMATODI 476  MAMMATODI 478  HAMMATODI 479  TAMMATODI 499  MAMMATODI 501  MAMMATODI 501  MAMMATODI 501  MAMMATODI 502  MAMMATODI 502  MAMMATODI 502  MAMMATODI 502  MAMMATODI 503	33. 639 117. 183 155. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000
	MAMATOD1476 MAMATOD1478 MAMATOD1478 MAMATOD1479 MAMATOD1498 MAMATOD1501 MAMATOD1501 MAMATOD1502 MAMATOD1510 MAMATOD1510 MAMATOD1522 MAMATOD1522 MAMATOD1522 MAMATOD1523 MAMATOD1532 MAMATOD1533	33. 639 117. 183 155. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537
	MAMATODI 476 MAMATODI 478 MAMATODI 479 MAMATODI 479 MAMATODI 489 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 502 MAMATODI 502 MAMATODI 502 MAMATODI 503 MAMATODI 503 MAMATODI 503 MAMATODI 503 MAMATODI 503 MAMATODI 503	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341	19. 551 61. 333 59. 931 51. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660
	MAMATODI 476  MAMATODI 478  MAMATODI 479  MAMATODI 487  MAMATODI 487  MAMATODI 501  MAMATODI 502  MAMATODI 502  MAMATODI 522  MAMATODI 522  MAMATODI 523  MAMATODI 532  MAMATODI 533  MAMATODI 534  MAMATODI 534  MAMATODI 535  MAMATODI 535  MAMATODI 535  MAMATODI 535  MAMATODI 535	33.639 117.183 156.131 67.613 96.522 216.969 124.674 27.993 56.601 83.190 47.058 97.390 0.341 32.482 122.717	19. 551 61. 333 59. 931 53. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0.000 23. 902 186. 325	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537
	MAMATOD 1476 MAMATOD 1478 MAMATOD 1479 MAMATOD 1479 MAMATOD 1487 MAMATOD 1501 MAMATOD 1502 MAMATOD 1502 MAMATOD 1502 MAMATOD 1522 MAMATOD 1522 MAMATOD 1522 MAMATOD 1523 MAMATOD 1532 MAMATOD 1532 MAMATOD 1534 MAMATOD 1534 MAMATOD 1534 MAMATOD 1535 MAMATOD 1534 MAMATOD 1535 MAMATOD 1535 MAMATOD 1535 MAMATOD 1535 MAMATOD 1535	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717	19. 551 61. 333 59. 931 53. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0.000 23. 902 186. 325 155. 615	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781
40	MAMATODI 476 MAMATODI 478 MAMATODI 478 MAMATODI 479 MAMATODI 497 MAMATODI 498 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 510 MAMATODI 510 MAMATODI 512 MAMATODI 522 MAMATODI 532 MAMATODI 533 MAMATODI 533 MAMATODI 534 MAMATODI 535 MAMATODI 537	33.639 117.183 156.131 67.613 96.522 216.969 124.674 27.993 56.601 83.190 47.058 97.390 0.341 32.482 122.717 103.124 47.916	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0.000 23. 902 186. 325 155. 615 56. 549	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 592 18. 319	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381
	MAMATODI 476 MAMATODI 478 MAMATODI 479 MAMATODI 479 MAMATODI 487 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 502 MAMATODI 510 MAMATODI 512 MAMATODI 522 MAMATODI 522 MAMATODI 532 MAMATODI 533 MAMATODI 534 MAMATODI 534 MAMATODI 536 MAMATODI 537 MAMATODI 537 MAMATODI 569 MAMMATODI 578	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156
40	MAMATODI 476 MAMATODI 478 MAMATODI 479 MAMATODI 479 MAMATODI 487 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 502 MAMATODI 502 MAMATODI 502 MAMATODI 503 MAMAT	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636
40	MAMATOD 1476 MAMATOD 1478 MAMATOD 1479 MAMATOD 1479 MAMATOD 1487 MAMATOD 1501 MAMATOD 1501 MAMATOD 1502 MAMATOD 1502 MAMATOD 1502 MAMATOD 1502 MAMATOD 1502 MAMATOD 1528 MAMATOD 1528 MAMATOD 1533 MAMATOD 1534 MAMATOD 1535 MAMATOD 1535 MAMATOD 1536 MAMATOD 1537 MAMATOD 1537 MAMATOD 1537 MAMATOD 1537 MAMATOD 1537 MAMATOD 1576	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860	19. 551 61. 333 59. 931 51. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 725 30. 090 57. 322 30. 398	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 45. 073 38. 692 18. 319 29. 095 62. 142 24. 245	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 50.896 115.580 30.161	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305
40	MAMATOD 1476  MAMATOD 1478  HAMMATOD 1479  CLAMATOD 1487  CLAMATOD 1487  MAMATOD 1501  MAMATOD 1501  MAMATOD 1502  MAMATOD 1502  MAMATOD 1502  MAMATOD 1522  MAMATOD 1522  MAMATOD 1523  MAMATOD 1532  MAMATOD 1534  MAMATOD 1534  MAMATOD 1535  MAMATOD 1536  MAMATOD 1537  MAMATOD 1537  MAMATOD 1537  MAMATOD 1536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157	19. 551 61. 333 59. 931 51. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 245 1. 210	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 580 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 11. 381 33. 156 39. 636 22. 305 4. 371
40	MAMATOD 1476  MAMATOD 1478  MAMATOD 1479  CLAMATOD 1479  CLAMATOD 1487  MAMATOD 1501  MAMATOD 1501  MAMATOD 1502  MAMATOD 1502  MAMATOD 1502  MAMATOD 1502  MAMATOD 1522  MAMATOD 1522  MAMATOD 1523  MAMATOD 1532  MAMATOD 1533  MAMATOD 1534  MAMATOD 1535  MAMATOD 1536  MAMATATOD 1569  MAMATATOD 1569  MAMATATOD 1566  MAMATOD 1566  MAMATATOD 1566  MAMATOD 1566  MAMATOD 1566  MAMATOD 1566  MAMATOD 1566  MAMATOD 1566	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157	19. 551 61. 333 59. 931 51. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.556 2.133 84.714	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869
40	MAMATOD 1476  MAMATOD 1478  MAMATOD 1479  MAMATOD 1479  MAMATOD 1487  MAMATOD 1501  MAMATOD 1501  MAMATOD 1502  MAMATOD 1502  MAMATOD 1522  MAMATOD 1522  MAMATOD 1523  MAMATOD 1523  MAMATOD 1524  MAMATOD 1525  MAMATOD 1534  MAMATOD 1535  MAMATOD 1535  MAMATOD 1535  MAMATOD 1536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616	19. 551 61. 333 59. 931 53. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889	25. 289 147. 393 31. 646 92. 480 222. 159 84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 45. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 37. 913 27. 880	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119
40	MAMATODI 476 MAMATODI 478 MAMATODI 478 MAMATODI 479 MAMATODI 497 MAMATODI 497 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 510 MAMATODI 510 MAMATODI 522 MAMATODI 522 MAMATODI 522 MAMATODI 523 MAMATODI 533 MAMATODI 533 MAMATODI 534 MAMATODI 535 MAMATODI 535 MAMATODI 535 MAMATODI 536 MAMATODI 537 MAMATODI 537 MAMATODI 537 MAMATODI 536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 889 32. 889 32. 647	25. 289 147 393 31. 646 92. 480 92. 480 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 45. 073 38. 592 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344
40 45	MAMATODI 476 MAMATODI 478 MAMATODI 479 MAMATODI 479 MAMATODI 487 MAMATODI 487 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 510 MAMATODI 510 MAMATODI 512 MAMATODI 522 MAMATODI 522 MAMATODI 533 MAMATODI 533 MAMATODI 534 MAMATODI 534 MAMATODI 535 MAMATODI 536 MAMAT	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112	19. 551 61. 333 59. 931 53. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 0. 000 21. 042 75. 842 52. 282 19. 725 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765	25. 289 147. 393 31. 646 92. 480 92. 480 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639 24.401 44.599	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253
40 45	MAMATODI 476 MAMATODI 478 MAMATODI 478 MAMATODI 479 MAMATODI 497 MAMATODI 497 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 510 MAMATODI 510 MAMATODI 522 MAMATODI 522 MAMATODI 522 MAMATODI 523 MAMATODI 533 MAMATODI 533 MAMATODI 534 MAMATODI 535 MAMATODI 535 MAMATODI 535 MAMATODI 536 MAMATODI 537 MAMATODI 537 MAMATODI 537 MAMATODI 536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616	19. 551 61. 333 59. 931 53. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 29. 88	25. 289 147. 393 31. 646 92. 480 92. 480 92. 480 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788	24. 584 32. 143 62. 901 40. 427 23. 385 88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253 78. 377
40 45	MAMATODI 476 MAMATODI 478 MAMATODI 478 MAMATODI 479 MAMATODI 487 MAMATODI 487 MAMATODI 501 MAMATODI 501 MAMATODI 502 MAMATODI 510 MAMATODI 510 MAMATODI 512 MAMATODI 522 MAMATODI 522 MAMATODI 533 MAMATODI 533 MAMATODI 534 MAMATODI 534 MAMATODI 535 MAMATODI 536 MAMAT	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112	19. 551 61. 333 59. 931 51. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469 23. 619	25. 289 147. 393 31. 646 92. 480 92. 480 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 45. 073 38. 692 29. 095 62. 142 24. 245 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639 24.401 44.599 15.339 88.514	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253 78. 377 13. 860
40 45	MAMMATODI 476 MAMMATODI 478 MAMMATODI 479 MAMMATODI 479 MAMMATODI 487 MAMMATODI 501 MAMMATODI 501 MAMMATODI 502 MAMMATODI 502 MAMMATODI 502 MAMMATODI 503 MAMMATODI 503 MAMMATODI 503 MAMMATODI 503 MAMMATODI 504 MAMMATODI 505 MAMMATODI 507 MAMMATODI 507 MAMMATODI 506 MAMMATODI 606 MAMMATODI 606 MAMMATODI 606 MAMMATODI 606 MAMMATODI 606	33. 539 117. 183 1156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088	19. 551 61. 333 59. 931 53. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 29. 88	25. 289 147. 393 31. 646 92. 480 92. 480 92. 480 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016 13.148 52.861	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639 24.401 44.599 15.339 88.514 9.100 35.992	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 580 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011 21. 716	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 13. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253 78. 377 13. 860 14. 483
40 45	MAMA 1001 476  MAMA 1001 478  HAMMA 1001 479  ELAMA 1001 487  HAMMA 1001 487  HAMMA 1001 487  HAMMA 1001 501  MAMA 1001 502  HAMMA 1001 502  HAMMA 1001 529  HAMMA 1001 529  HAMMA 1001 532  HAMMA 1001 534  HAMMA 1001 535  MAMMA 1001 535  MAMMA 1001 536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637 74. 839	19. 551 61. 333 59. 931 51. 042 111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469 23. 619	25. 289 147. 393 31. 646 92. 480 92. 480 92. 481 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919 74. 281	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016 13.148 52.861 91.848 18.302	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 45. 073 38. 692 29. 095 62. 142 24. 245 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639 24.401 44.599 15.339 88.514	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 33. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253 78. 377 13. 860
40 45	MAMA 1001 476  MAMA 1001 478  HAMMA 1001 479  ELAMA 1001 479  ELAMA 1001 487  ELAMA 1001 487  ELAMA 1001 501  ELAMA 1001 502  ELAMA 1001 502  ELAMA 1001 502  ELAMA 1001 522  ELAMA 1001 522  ELAMA 1001 523  ELAMA 1001 533  ELAMA 1001 534  ELAMA 1001 535  ELAMA 1001 536  ELAMA 1001 537  ELAMA 1001 536	33. 639 117. 183 156. 131 67. 613 96. 522 216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637	19. 551 61. 333 59. 931 51. 042 1111. 213 55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469 23. 619 29. 828	25. 289 147. 393 31. 646 92. 480 92. 480 92. 481 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919 74. 281 9. 202	5.909 46.785 28.808 34.978 50.813 38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000 24.788 45.519 43.540 24.376 31.981 39.259 23.526 2.133 84.714 14.016 13.148 52.861 91.848 18.302	17. 988 39. 649 44. 671 30. 928 14. 811 49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 245 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063 18. 016	24.584 32.143 62.901 40.427 23.385 88.169 54.854 6.993 26.994 41.170 23.522 25.298 6.274 27.839 43.338 47.685 34.666 50.896 115.580 30.161 6.758 67.639 24.401 44.599 15.339 88.514 9.100 35.992	26. 252 33. 776 69. 911 27. 489 56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 580 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011 21. 716	17. 981 40. 723 26. 759 11. 238 28. 054 32. 036 36. 860 8. 048 17. 416 20. 596 30. 896 16. 233 0. 000 10. 537 24. 660 32. 781 11. 381 13. 156 39. 636 22. 305 4. 371 52. 869 19. 119 20. 344 16. 253 78. 377 13. 860 14. 483

Table 56

				22 183 1	40 00 1	177 660 1	125 761 T	44 200
MARYA1001619	361.714	66.104	138, 945	35. 137	88.004	177. 280	155.721	44. 365
MALMA1001620	113. 233	68.799	320.014	88. 182	65. 387	62.891	47.797	49. 428
MALMA1001623	32.719	16.493	22.246	8.396	13.561	16. 233	7, 490	7.940
MALMA1001626	75. 279	8.514	13.728	10.774	12.665	56.613	57.493	6.962
4AP4A1001627	28. 468	7.652	39. 356	8.734	4.064	8. 190	14. 443	7.576
MAYMA1001630	36.419	36.649	115. 287	20. 971	7.371	8.511	10.371	16.570
HAPMA1001633	77.945	25. 597	143.786	22.273	51.279	40.689	37. 952	19. 350
MAPMA1001634	132, 937	95, 570	297.140	83.974	56.835	62.263	58. 952	66.333
MALYA1001635	140.754	47.359	225. 161	34. 126	24,717	38.086	34.792	34.698
MALMA1001649	30. 569	12.321	20.513	11.727	13,713	19, 299	12.550	9.106
HAPMA1001654	150, 282	91.691	90.096	34.969	64. 959	66.853	62.712	58. 197
MALMA1001660	133.470	97.805	42.199	61.020	54.089	65.813	66.019	54. 874
MALMA1001563	394.964	202.523		154. 372	162.177	148.843	118. 542	79. 262
MAPMA1001670	109.171	38. 230	119.077	31.362	18.030	43.797	53.194	28. 426
HALMA1001671	145. 809	21.188	31.621	20. 983	11.973	13.009	10.867	8.816
HAE M1001679	74. 490	17.313	20. 426	10. 837	8.375	23. 180	9. 271	18.786
			260.375	71.605	39.630	48. 331	49. 633	41.012
MALMA1001683	147.044	87.078	46.223	12.860	21.575	12. 528	5.274	9.906
MALMA1001686	12.824	14. 464		407. 762	105.060	319.616		1824. 587
MALMA1001688	290.960	584.756 1					31.387	18. 923
MAPMA1001689	74.686	28. 294	39.725	20.248	8. 261	19.721	16.899	28. 825
MAPMA1001692	90. 375	64.474	198.053	56. 976	35. 470	19.914	10.898	27.229
MAPMA1001711	111. 425	82.300	189. 195	30. 269	36.663	51.227		13. 907
MAKMA1001715	67.545	40. 330	71.553	28.616	19.372	25.019	24. 223	8. 587
MAPMA1001730	33.925	17.096	21.837	11.464	4, 477	36.743	11.375	
MAMMA1001735	79.384	42.172	38. 240	23.675	25. 390	20.932	27.963	11.313
MANMA 1001740	100.894	25. 218	94. 454	17.836.	17.794	23.366	21.945	16. 107
MARSA 1001743	199, 112	118, 364	141.535	72.049	46. 384	86.104	96.828	100.038
HALEMA1001744	23. 256	20. 454	0.000	2.086	2, 551	2.098	5.703	0.000
MAK≅A1001745	121.679	94. 047	301.292	106. 455	100, 677	125.697	46.388	55.894
MAMMA 1001751	58.670	37. 967	90.572	30. 921	14.618	26.060	33.416	32. 380
MAMMA 100 1752	284. 221	89.024	175.680	74.746	86.008	159.864	103.908	99.685
MANMA1001754	57.620	30. 193	53. 390	14. 833	35. 182	39.454	17.523	12.754
MANMA1001757	14. 456	8. 290	7.632	7.247	6.076	15.580	5. 382	5.641
MAMMA 1001760	283. 527	155. 103	596.815	118. 229	106.868	115.717	105. 154	147.707
MANNA 1001764	33.825	15.661	31.885	14. 429	5.043	11.697	22.420	16. 539
MANNA 1001767	41.791	27.578	112. 242	22. 484	21.848	16.357	11.576	9. 367
MALSA 1001768	50,861	34.645	129.707	25. 692	23.037	24.674	27.811	11.075
MALMA1001769	206.737	82.818	645.195	110.913	102.640	105.607	80.653	102.144
HAMMA1001771	123.973	30.551	49.772	16.877	55.099	52.348	41.113	48.806
MAMMA 1001773	47.743	27.204	35. 277	8. 450	18.002	17, 141	23.713	30.755
MALMA 1001778	104.585	49.619	92.589	42. 249	35.085	50.584	39.215	26.862
MAMMA 1001783	140.821	89.274	371.095	82. 231	85.003	87.248	61.999	71.448
MANMA 1001785	119.072	65.819	256.400	60. 491	37.351	65.802	45.875	54.652
MANMATO01788	37.967	8.305	25. 708	9.749	9.870	11.494	13.172	10.408
MANMA1001790	202.092	181. 258	279. 482	57, 700	22.737	29. 284	28.819	46.106
MANAA 1001800	24. 282	11.444	30.466	12.517	1.763	8.501	13.065	25. 671
MANSAA 100 1804	150.744	16.771	51.213	14. 975	33.630	67.533	64.799	20.701
HAMMA1001806	62.312	54. 896	146, 142	37.371	11.402	36.501	43.675	52.846
MAMMA 1001812	17.002	11.569	32.023	10.166	5. 995	9.576	10.245	11.255
MAMMA 1001815	50.743	27, 272	61.778	19.704	15. 636	25.863	15. 187	22.130
MALSMA 1001817	10.653	7.578	15.446	7.044	7.758	3.511	7.974	11.601
HAMMA1001818	48.733	19.657	87. 193	21.647	18.566	18.770	19. 255	18.678
MAN=4A1001819	165. 340	99. 233	343, 318	111.523	112.261	57.848	73, 268	87.725
MAPSA 1001820	48.662	22.951	34, 879	15. 243	11.743	9.468	15.897	11.396
MAMMA1001824	125. 683	53.824	187. 383	58.214	53, 691	47.999	45. 347	37. 548
		30.370	42.082	21.957	23.518	23.996	20.046	8, 482
	1 36 644	1 00.010		45.913	43.465	44.952	56.814	24.346
MANMA 1001832	56.633	58 280	1 1/4 NA!					50.30
MAMMA 1001832 MARMA 1001836	128.477	58. 280	179.541		38 152	37 090	17.947	
MAMMA1001832 MAMMA1001836 MAMMA1001837	128. 477	65.031	172.658	60.299	38.153	37.090	17.947	
MAMMA 1001832 MAMMA 1001836 MAMMA 1001837 MAMMA 100183U	128. 477 118. 428 42. 562	66.031 27.622	172.658 82.759	60. 299 24. 693	20.435	22.941	15. 102	19. 124
MAMMA 1001832 MAMMA 1001836 MAMMA 1001837 MAMMA 1001834 MAMMA 1001850	128. 477 118. 428 42. 562 402. 506	66. 031 27. 622 243. 182	172.658 82.759 312.586	60.299 24.693 171.182	20. 435 143. 034	22.941 232.615	15. 102 91. 466	19.124 106.637
MAMMA 1001832 MAMMA 1001836 MAMMA 1001837 MAMMA 1001840 MAMMA 1001850 MAMMA 1001851	128. 477 118. 428 42. 562 402. 506 123. 305	66. 031 27. 622 243. 182 30. 035	172.658 82.759 312.586 69.870	60. 299 24. 693 171. 182 64. 763	20. 435 143. 034 41. 560	22.941 232.615 39.454	15. 102 91. 466 33. 329	19. 124 106. 637 45. 924
MAMMA 1001832 MAMMA 1001836 MAMMA 1001837 MAMMA 1001848 MAMMA 1001850 MAMMA 1001851 MAMMA 1001852	128. 477 118. 428 42. 562 402. 506 123. 305 198. 774	66.031 27.622 243.182 30.035 161.311	172.658 82.759 312.586 69.870 321.896	60. 299 24. 693 171. 182 64. 763 118. 228	20. 435 143. 034 41. 560 133. 655	22.941 232.615 39.454 112.820	15.102 91.466 33.329 91.724	19. 124 106. 637 45. 924 115. 602
MAMMA 1001832 MAMMA 1001836 MAMMA 1001837 MAMMA 1001840 MAMMA 1001850 MAMMA 1001851	128. 477 118. 428 42. 562 402. 506 123. 305	66. 031 27. 622 243. 182 30. 035	172.658 82.759 312.586 69.870	60. 299 24. 693 171. 182 64. 763	20. 435 143. 034 41. 560	22.941 232.615 39.454	15. 102 91. 466 33. 329	19. 124 106. 637 45. 924 115. 602 45. 321 58. 279

Table 57

BMCSAIDORS64   169, 742   52,389   185,785   37,880   50,386   67,995   55,272   23,142										
5		HAKMA1001864	169.742	52.389	185. 785	37. 880	50.896	67.999	55. 272	23, 142
### RACKAIODIAPS 190. 515		MALEMA 1001868	82.643	56.439	59. 491	62.418	34. 438	47.003	29.588	_35.585
Section   Color   Co		MALYA1001874	9, 192	9,651	51, 178	7, 405	11.275	9.054	7.189	10.453
MACKANDO	5						101, 886	72, 219	79.645	146, 982
NUMPSINGO 1895   17, 729   44, 975   10, 556   55, 546   26, 142   52, 223   41, 423   29, 156										
NAMES   100   1890   177   595   47, 712   247, 584   60, 586   29, 367   36, 338   39, 109   41, 483										
TAMPSINGO 1993   30, 120   22, 271   50, 435   19, 070   22, 222   27, 783   36, 543   18, 711										
The color of the										
DEMENTOD1907   159.767   70.062   305.846   76.004   91.563   25.690   68.288   28.95										
### Comparison   44.5   44.7   77.928   47.5   57.352   40.219   53.908   32.123   40.375   ### Comparison   34.5   57.5   58.8   37.5			78.854	67.274	188.894					
### HEXAGIODISTS 9.659 2.713 43.522 17.305 10.000 2.270 0.000 0.000 1.000 1.051 15.715 184241001931 5.716 9.659 2.713 43.522 13.591 18.155 29.466 11.457 184241001937 14.7045 76.457 31.002 16.635 17.844 31.265 29.899 19.500 10.24101935 14.7045 76.465 31.1021 16.635 17.844 31.265 29.899 19.500 10.24101935 14.304 40.783 17.518 29.530 76.771 16.554 47.425 67.568 75.411 19.241101936 1714.304 40.783 15.55.368 46.619 41.439 44.871 67.553 25.7411 19.241101936 19.822 63.449 192.955 55.222 57.938 21.395 42.027 43.871 63.25 29.899 19.24 20.27 41.871 60.2854 19.241101936 19.822 13.449 192.955 55.222 57.938 21.395 42.027 42.871 62.844 19.241101936 19.822 13.449 192.955 55.222 57.938 21.395 42.027 42.871 62.844 19.241101936 19.822 13.449 192.955 55.222 57.938 21.395 42.027 42.871 62.844 19.241101936 19.823 10.	10	MARSUA 1001907	159.767		305. 846					
The color of the		MALMA 1001908	44.964	27.928	41.967	55, 852	40.219	53.008		
SECTION   STATE   ST		9191001919	0.000	82,865	12, 109	0.000	2.270	0.000	0.000	5. 175
BACCANDOISST   14.7045   75.453   33.102   16.535   17.844   31.265   29.899   19.650								18. 165	29.466	11.467
HARMANDOTSES   17.033   17.5   574   31.618   70.511   55.661   40.552   39.990   40.224								31, 265	29,899	19.650
### PREMAINDUSTS   171, 199   78, 116   295, 300   76, 171   65, 654   47, 426   67, 568   57, 411   ### PREMAINDUSTS   11, 104   40, 789   155, 156   46, 819   41, 429   43, 671   20, 153   52, 615   32, 692   ### PREMAINDUSTS   13, 104   40, 789   155, 156   46, 819   41, 429   43, 671   20, 33   35   42, 027   44, 844   ### PREMAINDUSTS   15, 313   1, 651   3, 748   3, 671   3, 337   0, 000   0, 000   0, 000   5, 775   ### PREMAINDUSTS   17, 705   0, 000   0, 000   0, 000   1, 158   41, 691   71, 805   71, 805   61, 125   ### PREMAINDUSTS   17, 705   0, 000   0, 000   0, 000   0, 000   1, 681   1, 561   0, 000   0, 000   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   283, 440   78, 807   70, 640   63, 218   71, 282   22, 888   ### PREMAINDUSTS   189, 502   91, 530   143, 529   143, 289   143, 289   ### PREMAINDUSTS   189, 502   91, 530   143, 529   173, 534   53, 243   47, 173   47, 183   ### PREMAINDUSTS   189, 502   91, 530   189, 530   10, 538   10, 384   48, 48, 48, 48, 48, 48, 48, 48, 48, 48,										
MACHING   1978   114, 304   40, 789   155, 365   66, 819   41, 429   43, 871   26, 153   36, 982   144, 1424   142, 143   142, 143   143, 143, 143   143, 143, 143, 143, 143, 143, 143, 143,	15									
MACHAN HOOT SEG	13									
Characteristics   Characteri										
### DRESS   103   159   154   159   157   158   149   150   109   284   97   612   137   120   ### DRESS   159   159   159   158   123   123   1										
Colorador   199   358   123   085   297   080   01   158   41   691   71   806   71   681   61   125										
Color										
PAMEMIO   1997   189   507   91   630   283   440   78   807   70   640   63   218   71   282   32   388   144   140		MARMA 1001970								
EMPERATION   Section   S	20	MAMMA1001978	1.206	0.000						
EMMERIO02008   66.834   77.793   37.647   14.813   20.016   33.334   39.965   10.388   EMMERIO02009   144.462   65.030   407.311   107.350   55.438   47.107   40.434   57.138   EMMERIO02019   144.462   65.030   407.311   107.350   55.438   47.107   40.434   57.138   EMMERIO02022   107.727   67.057   159.576   65.640   59.239   37.381   36.122   50.747   50.74		MAMMA1001992	189. 502	91.630	283.440	78.807	70.640	63.218	71.282	
EMBEST 002009		MAMMA1001994	85. 231	21.385	143. 259	40. 178	38. 484	54. 686	24.893	33.837
MARINATORZO09   144, 452   65, 030   407, 911   107, 350   55, 438   47, 107   40, 434   57, 138     MARINATORZO11   32, 832   13, 901   27, 624   10, 188   19, 701   17, 344   22, 354   44, 449     MARINATORZO22   176, 885   70, 125   207, 390   72, 614   55, 279   78, 953   108, 945   46, 948     MARINATORZO22   270, 523   310, 983   362, 313   98, 620   95, 523   104, 970   73, 966   83, 780     MARINATORZO23   176, 885   70, 125   207, 390   72, 614   55, 279   78, 953   108, 945   46, 948     MARINATORZO24   176, 885   70, 125   207, 390   72, 614   55, 279   78, 953   108, 945   46, 948     MARINATORZO24   19, 611   15, 313   18, 901   14, 070   10, 859   15, 705   11, 098   10, 476     MARINATORZO24   78, 700   42, 958   161, 397   37, 586   30, 208   55, 485   24, 562   23, 890     MARINATORZO25   71, 31   8, 948   24, 018   14, 459   14, 811   11, 172   1, 533   10, 371     MARINATORZO265   212, 189   152, 223   474, 785   146, 238   94, 517   84, 218   104, 806   75, 923     MARINATORZO265   212, 189   152, 223   474, 785   146, 238   94, 517   84, 218   104, 806   75, 923     MARINATORZO265   128, 185   46, 405   127, 810   82, 855   59, 107   72, 737   65, 052   39, 657     MARINATORZO265   128, 185   46, 405   127, 810   82, 855   59, 107   72, 737   65, 052   39, 657     MARINATORZO265   128, 185   46, 405   127, 810   82, 855   59, 107   72, 737   65, 052   39, 657     MARINATORZO265   128, 186   24, 791   29, 988   16, 102   15, 306   31, 362   22, 002   21, 338     MARINATORZO265   128, 186   24, 791   29, 988   16, 102   15, 306   31, 362   22, 002   21, 338     MARINATORZO267   17, 991   17, 619   30, 479   8, 932   13, 026   13, 362   22, 002   21, 338     MARINATORZO267   17, 991   17, 619   30, 479   8, 932   13, 026   13, 365   9, 996   63, 344     MARINATORZO267   17, 991   17, 619   30, 479   8, 932   13, 026   13, 365   9, 996   63, 344     MARINATORZO267   17, 991   17, 619   30, 479   8, 932   31, 302   31, 362   22, 002   21, 388     MARINATORZO268   158, 168   24, 279   27, 377		MAMMA1002008	66.834	77.793	37, 647	14.813	20.016	33. 334	39. 365	10.388
EMEMAT002011   32,832   13,901   27,624   10,188   19,701   17,344   22,354   14,449							55. 438	47.107	40.434	57.138
CAMBIAT 1002022   107.727   67.057   159.576   65.640   59.239   37.381   36.122   50.747								17, 344	22, 354	14.449
CAMBAI 1002024									36, 122	50.747
Figure   1002032   270.523   130.983   562.313   98.620   95.826   104.970   73.966   83.780	25									
日本語本1002033										
### PART   19,611   15,313   18,901   14,070   10,859   15,705   11,098   10,476										
######################################										
SAME										
CAMENAT 100 2047   82.875   57.343   192.240   55.806   45.781   34.315   27.824   37.210     MANIENAT 100 2056   212.189   152.323   474.785   146.238   94.617   84.218   104.806   75.923     CAMENAT 100 2056   149.112   126.148   334.116   98.541   74.809   81.670   44.227   55.825     MANIENAT 100 2058   149.112   126.148   334.116   98.541   74.809   81.670   44.227   55.825     MANIENAT 100 2058   128.185   46.405   127.810   82.855   59.107   72.737   63.052   39.667     MANIENAT 100 2058   110.652   64.982   163.753   51.583   45.893   40.656   37.400   24.128     MANIENAT 100 2070   61.186   24.791   29.988   16.102   15.306   31.362   22.002   21.338     MANIENAT 100 2070   170.197   38.633   93.014   30.633   33.682   90.533   42.110   14.299     MANIENAT 100 2080   21.195   14.596   12.646   10.208   14.094   14.792   10.377   10.263     MANIENAT 100 2084   74.297   40.086   152.790   30.118   30.052   28.788   24.428   24.140     MANIENAT 100 2084   74.297   40.086   152.790   30.118   30.052   28.788   24.428   24.140     MANIENAT 100 2083   77.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     MANIENAT 100 2085   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     MANIENAT 100 2019   77.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     MANIENAT 100 2019   77.898   6.035   31.027   13.639   7.393   32.486   27.923   11.735     MANIENAT 100 2119   122.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     MANIENAT 100 2119   122.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     MANIENAT 100 2112   24.376   27.337   10.667   11.574   5.250   15.678   14.329   37.465     MANIENAT 100 2112   22.675   118.230   198.712   79.589   88.860   84.266   50.530   48.550     MANIENAT 100 2112   21.646   33.612   49.214   19.085   27.295   103.698   63.341   39.193   25.406     MANIENAT 100 2142   21.646   33.612   49.214   19.085   27.295   103.698   63.348   39.519     MANIENAT 100 2142   21.646   33.612   49.214   19.085										
Main	30									
CAMMA1002058   143.112   126.148   334.116   98.541   74.809   81.670   44.227   65.825     CAMMA1002060   13.278   7.931   14.514   12.643   5.782   6.917   16.902   5.536     MAMMA1002065   128.185   46.405   127.810   82.855   59.107   72.737   63.052   39.667     CAMMA1002068   110.652   64.982   163.753   51.583   45.893   40.656   37.400   24.128     CAMMA1002070   61.186   24.791   29.988   16.102   15.306   31.362   22.002   21.338     CAMMA1002070   170.197   38.633   93.014   30.633   33.682   90.533   42.110   14.299     CAMMA1002002   111.870   77.716   117.819   55.009   54.940   28.457   25.946   21.254     CAMMA1002004   74.297   40.086   152.790   30.118   30.052   28.788   24.428   24.140     CAMMA1002007   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CAMMA1002007   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CAMMA1002007   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CAMMA1002007   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CAMMA1002007   17.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     CAMMA1002007   17.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     CAMMA1002007   17.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     CAMMA1002107   12.2271   36.908   59.513   20.581   36.895   32.486   27.923   11.735     CAMMA1002112   24.376   27.337   10.667   11.574   5.250   15.678   4.329   37.463     CAMMA1002112   12.271   36.908   59.513   20.581   36.895   38.172   39.466   32.476     CAMMA1002112   12.271   36.908   59.513   20.581   36.895   38.172   39.463   32.476     CAMMA1002126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     CAMMA1002126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     CAMMA1002126   231.380   33.612   49.214   19.085   27.295   103.698   68.348   39.850     CAMMA1002127   21.646   33.612   49.214   19.085   27.295   103.698   68.348   39.850     CAMMA100	50									
CLAMMATORZOBS   12.8.185   46.405   127.810   82.855   59.107   72.737   63.052   39.667   13.061   10.0652   12.8.185   46.405   127.810   82.855   59.107   72.737   63.052   39.667   13.061   10.0652		MAISMA1002056	212. 189							
HAMMA1002085   128.185   46.405   127.810   82.855   59.107   72.737   63.052   39.667     CASMA1002088   110.652   64.982   163.753   51.583   45.893   40.656   37.400   24.128     CASMA1002070   61.186   24.791   29.988   16.102   15.306   31.362   22.002   21.338     CASMA1002088   170.197   38.633   93.014   30.633   33.682   90.533   42.110   14.299     CASMA1002080   21.195   14.596   12.646   10.208   14.094   14.792   10.377   10.263     CASMA1002084   74.297   40.086   12.646   10.208   14.094   14.792   10.377   10.263     CASMA1002087   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CASMA1002097   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CASMA1002091   78.604   26.611   41.258   17.086   26.812   39.757   46.803   27.660     CASMA1002093   77.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     CASMA1002095   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     CASMA1002012   24.376   27.337   10.667   11.574   5.250   15.678   14.329   37.463     CASMA1002112   24.376   27.337   10.667   11.574   5.250   15.678   14.329   37.463     CASMA1002113   12.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     CASMA1002114   12.060   5.100   8.756   5.943   6.502   7.856   7.396   3.149     CASMA1002115   122.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     CASMA1002126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     CASMA1002126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     CASMA1002128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     CASMA1002128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     CASMA1002128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     CASMA1002143   150.595   15.368   78.681   38.118   5.385   13.974   10.806   45.937     CASMA1002143   150.595   15.368   78.681   38.118   5.385   13.974   10.806   45.937		MANMA1002058			334.116	98. 541				
TANSA   1002068   10.652   64.982   163.753   51.583   45.893   40.656   37.400   24.128		MAMMA1002060	13. 278	7, 931	14.514	12.643				
TANSATOO21970   61.186   24.791   29.988   16.102   15.306   31.362   22.002   21.338     TANSATOO2078   170.197   38.633   93.014   30.633   33.682   90.533   42.110   14.299     TANSATOO2080   21.195   14.596   12.646   10.208   14.094   14.792   10.377   10.263     TANSATOO2082   111.870   77.716   117.819   55.009   54.940   28.457   25.946   21.254     TANSATOO2087   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     TANSATOO2087   17.991   17.619   30.479   8.932   13.026   13.365   9.996   6.344     TANSATOO2087   17.498   0.000   5.942   5.592   5.630   8.103   11.278   4.689     TANSATOO2085   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     TANSATOO2085   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     TANSATOO2108   91.919   6.035   31.027   13.639   7.939   32.486   27.923   11.735     TANSATOO2118   12.060   5.100   8.756   5.943   6.502   7.856   7.396   31.49     TANSATOO2118   12.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     TANSATOO2118   12.277   33.844   373.786   60.523   54.991   63.367   35.366   35.797     TANSATOO2126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     TANSATOO2128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     TANSATOO2140   54.642   53.277   115.593   42.121   33.524   31.026   24.905   32.121     TANSATOO2143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     TANSATOO2143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     TANSATOO2143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     TANSATOO2147   73.366   34.088   45.076   77.984   33.648   53.571   33.082   8.766     TANSATOO2147   73.366   34.088   45.076   77.984   33.648   53.571   33.082   8.766     TANSATOO2147   73.366   34.088   45.076   77.984   33.648   53.571   33.082   8.766     TANSATOO2147   73.366   34.088   45.076   77.984   33.648   53.571   33.082   8.766     TANSATOO2143		MAMMA1002065	128. 185	46. 405		82.855	59. 107	72.737	63.052	
CAMERIA1002070   61.186   24.791   29.988   16.102   15.306   31.362   22.002   21.338     CAMERIA1002080   170.197   38.633   93.014   30.633   33.682   90.533   42.110   14.299     CAMERIA1002080   21.195   14.596   12.646   10.208   14.094   14.792   10.377   10.263     CAMERIA1002084   74.297   40.086   152.790   30.118   30.052   28.788   24.428   24.140     CAMERIA1002097   77.916   17.619   30.479   8.932   13.026   13.365   9.996   6.344     CAMERIA1002097   78.604   26.611   41.258   17.086   26.812   39.757   46.803   27.660     CAMERIA1002091   78.604   26.611   41.258   17.086   26.812   39.757   46.803   27.660     CAMERIA1002095   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     CAMERIA1002096   78.790   13.430   22.728   13.058   20.650   32.157   32.621   8.152     CAMERIA1002108   91.919   6.035   31.027   13.639   7.939   32.486   27.923   11.735     CAMERIA1002112   24.376   27.337   10.667   11.574   5.250   15.678   14.329   37.463     CAMERIA1002119   122.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     CAMERIA1002119   122.271   36.908   59.513   20.581   36.895   38.172   39.046   32.476     CAMERIA1002128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     CAMERIA1002132   226.752   118.230   198.712   79.589   88.860   84.266   50.630   48.550     CAMERIA1002142   21.646   33.612   49.214   19.085   27.295   103.698   68.348   39.350     CAMERIA1002143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     CAMERIA1002143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     CAMERIA1002147   73.366   34.088   45.076   27.984   33.548   53.571   33.082   8.766     CAMERIA1002147   73.366   34.088   45.076   27.984   33.548   53.571   33.082   8.766     CAMERIA1002147   73.366   34.088   45.076   27.984   33.548   53.571   33.082   8.766     CAMERIA1002147   73.366   34.088   45.076   27.984   33.548   53.571   33.082   8.766     CAMERIA1002147   73.366   34.088   45.076   2		MAISMA1002068	110.652	64. 982	163.753	51.583	45.893	40.656		24. 128
CAMPAIA 1002008	35	MANSA 1002070		24, 791		16.102	15. 306	31.362	22.002	21.338
CLARMA1002080   21.195   14.596   12.646   10.208   14.094   14.792   10.377   10.263								90, 533	42.110	14.299
MAMMA1002092									10,377	10.263
### ### ##############################									25, 946	21, 254
### 17.619   30.479   8.932   13.026   13.365   9.996   6.344   ###################################										
######################################										
MAMATOOZO93   17,498   0,000   5,942   5,592   5,630   8,103   11,278   4,689     MAMATOOZO95   78,790   13,430   22,728   13,058   20,650   32,157   32,621   8,152     MAMATOOZO95   78,790   13,430   22,728   13,058   20,650   32,157   32,621   8,152     MAMATOOZO16   91,919   6,035   31,027   13,639   7,939   32,486   27,923   11,735     MAMATOOZO17   24,376   27,337   10,667   11,574   5,250   15,678   14,329   37,463     MAMATOOZO18   122,271   36,908   59,513   20,581   36,895   38,172   39,046   32,476     MAMATOOZO19   122,271   36,908   59,513   20,581   36,895   38,172   39,046   32,476     MAMATOOZO15   159,277   83,844   373,786   60,523   54,991   63,367   35,366   35,797     MAMATOOZO16   231,380   139,298   431,047   153,496   117,027   84,728   70,558   62,381     MAMATOOZO16   231,380   39,298   431,047   153,496   117,027   84,728   70,558   62,381     MAMATOOZO17   126,5752   118,230   198,712   79,589   88,860   84,266   50,630   48,550     MAMATOOZO140   54,642   53,227   115,593   42,121   33,524   31,026   24,905   32,121     MAMATOOZO142   121,646   33,612   49,214   19,085   27,295   103,698   68,348   39,836     MAMATOOZO143   150,595   15,368   78,681   38,118   5,895   13,974   10,806   45,937     MAMATOOZO147   73,366   34,088   45,076   27,984   33,648   53,571   33,082   8,766     MAMATOOZO153   133,485   74,073   143,431   55,132   46,673   85,911   25,126   19,099	40									
######################################	40									
TAPMA1002108										
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######################################										
HAMMA1002119 122.271 36.908 59.513 20.581 36.895 38.172 39.046 32.476 HAMMA1002125 159.277 83.844 373.786 60.523 54.991 63.367 35.366 35.797 HAMMA1002126 231.380 139.298 431.047 153.496 117.027 84.728 70.558 62.381 HAMMA1002132 102.647 35.864 48.863 19.098 20.911 44.235 39.193 25.406 HAMMA1002132 226.752 118.230 198.712 79.589 88.860 84.266 50.630 48.550  HAMMA1002140 54.642 53.227 115.593 42.121 33.524 31.026 24.905 32.121  MAMMA1002142 121.646 33.612 49.214 19.085 27.295 103.698 68.348 39.850  HAMMA1002143 150.595 15.368 78.681 38.118 5.895 13.974 10.806 45.937  HAMMA1002145 237.202 72.397 165.166 45.537 53.986 87.872 73.605 22.437  HAMMA1002147 73.366 34.088 45.076 27.984 33.648 53.571 33.082 8.766										<del></del>
HAMMA1002125   159.277   83.844   373.786   60.523   54.991   63.367   35.366   35.797     MAMMA1002126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     MAMMA1002128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     MAMMA1002132   226.752   118.230   198.712   79.589   88.860   84.266   50.630   48.550     MAMMA1002140   54.642   53.227   115.593   42.121   33.524   31.026   24.905   32.121     MAMMA1002142   121.646   33.612   49.214   19.085   27.295   103.698   68.348   39.850     MAMMA1002143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     MAMMA1002147   237.202   72.397   165.166   45.537   53.986   87.872   73.605   22.437     MAMMA1002147   73.366   34.088   45.076   27.984   33.648   53.571   33.082   8.766     MAMMA1002153   133.485   74.073   143.431   55.132   46.673   85.911   25.126   19.099										
INAMA   1002  125   159.277   83.844   373.786   60.523   54.991   63.367   35.366   35.797     INAMA   1002  126   231.380   139.298   431.047   153.496   117.027   84.728   70.558   62.381     INAMA   1002  128   102.647   35.864   48.863   19.098   20.911   44.235   39.193   25.406     INAMA   1002  132   226.752   118.230   198.712   79.589   88.860   84.266   50.530   48.550     INAMA   1002  140   54.642   53.227   115.593   42.121   33.524   31.026   24.905   32.121     INAMA   1002  142   121.646   33.612   49.214   19.085   27.295   103.698   68.348   39.850     INAMA   1002  143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     INAMA   1002  145   237.202   72.397   165.166   45.537   53.986   87.872   73.605   22.437     INAMA   1002  147   73.366   34.088   45.076   27.984   33.648   53.571   33.082   8.766     INAMA   1002  137   133.485   74.073   143.431   55.132   46.673   85.911   25.126   19.099	45									
#####################################			159. 277							
HAMMA1002132   226.752   118.230   198.712   79.589   88.860   84.266   50.630   48.550     MAMMA1002140   54.642   53.227   115.593   42.121   33.524   31.026   24.905   32.121     MAMMA1002142   121.646   33.612   49.214   19.085   27.295   103.698   68.348   39.850     MAMMA1002143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     MAMMA1002145   237.202   72.397   165.166   45.537   53.986   87.872   73.605   22.437     MAMMA1002147   73.366   34.088   45.076   27.984   33.648   53.571   33.082   8.766     MAMMA1002153   133.485   74.073   143.431   55.132   46.673   85.911   25.126   19.099		MAPMA1002126	231.380	139. 298	431.047	153.496	117.027			
MAMA1002132   226.752   118.230   198.712   79.589   88.860   84.266   50.630   48.550     MAMA1002140   54.642   53.227   115.593   42.121   33.524   31.026   24.905   32.121     MAMA1002142   121.646   33.612   49.214   19.085   27.295   103.698   68.348   39.825     MAMA1002143   150.595   15.368   78.681   38.118   5.895   13.974   10.806   45.937     MAMA1002143   237.202   72.397   165.166   45.537   53.986   87.872   73.605   22.437     MAMA1002147   73.366   34.088   45.076   27.984   33.648   53.571   33.082   8.766     MAMA1002153   133.485   74.073   143.431   55.132   46.673   85.911   25.126   19.099		MAMMA 1002128	102.647	35, 864	48.863	19.098	20. 911			
50 MA™A1002140 54.642 53.227 115.593 42.121 33.524 31.026 24.905 32.121 MA™A1002142 121.646 33.612 49.214 19.085 27.295 103.698 68.348 39.850 MA™A1002143 150.595 15.368 78.681 38.118 5.895 13.974 10.806 45.937 MA™A1002145 237.202 72.397 165.166 45.537 53.986 87.872 73.605 22.437 MA™A1002147 73.366 34.088 45.076 27.984 33.648 53.571 33.082 8.766 MA™A1002153 133.485 74.073 143.431 55.132 46.673 85.911 25.126 19.099		MAMMA1002132		118, 230	198.712	79.589	88.860	84. 266	50.630	48.550
50 NAMMA1002142 121.646 33.612 49.214 19.085 27.295 103.698 68.348 39.850								31.026	24, 905	32121
50 TIANMATODZI43 150.595 15.368 78.681 38.118 5.895 13.974 10.806 45.937 PIANMATODZI45 237.202 72.397 165.166 45.537 53.986 87.872 73.605 22.437 PIANMATODZI4Y 73.366 34.088 45.076 27.984 33.648 53.571 33.082 8.766 PIANMATODZI53 133.485 74.073 143.431 55.132 46.673 85.911 25.126 19.099									68.348	39.850
HAPMA1002145 237.202 72.397 165.166 45.537 53.986 87.872 73.605 22.437 HAPMA1002147 73.366 34.088 45.076 27.984 33.648 53.571 33.082 8.766 HAPMA1002153 133.485 74.073 143.431 55.132 46.673 85.911 25.126 19.099	50									
HAPMA1002147 73.366 34.088 45.076 27.984 33.648 53.571 33.082 8.766 HAPMA1002153 133.485 74.073 143.431 55.132 46.673 85.911 25.126 19.099										
TIAL A1002153 133.485 74.073 143.431 55.132 46.673 85.911 25.126 19.099										
130.400 100.400 14.010 140.401 40.102										
THE THIUM (155   120, 181   140, 215   332, 131   80, 240   120, 814   124, 336   33, 183   30, 376						66 44A				
		TAC 34 1002 155	1 320. 181	1 140. 215	1 224. 191	1 00. 240	1 120. 814	1 124. 330	33. 103	1 30.010

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Table 58

			Iai	)(C )(C			_	
HAMMA1002156	3, 612	2.088	14.013	0.880	0.000	0.000	0.000	0.554
HAMMA1002158	70. 916	40, 655	88.575	24, 125	14.786	21, 144	21.721	31.526
HAMMA1002164	109, 211	29. 584	54. 163	32.089	28.633	66.844	29.378	23.138
			135. 468	73.710	66.970	77.137	88. 540	53, 125
HALMA1002165	156.029	111.787			0.000	0.000	0.000	0.000
HAMMA 1002170	0.000	0.000	0.000	1, 159				
HALEMA 1002174	139. 902	178. 299		182. 252	147.225	141. 399	87.695	71.650
MANYA1002175	49.635	20.661	21.290	16.108	13, 918	22.449	12.876	18.564
HAMMA1002180	117.470	55.089	69.154	18.969	36.764	45. 946	59.721	45. 237
MAPMA1002198	123. 227	67. 539	235.488	54.699	51.835	48.796	31. 324	62.413
HALMA1002205	114.861	63, 437	420.688	47.331	61,775	61.499	42.296	74.029
HAPMA1002206	86. 539	30, 665	50.318	17.788	32.139	63. 320	64.272	56.392
HALMA1002209	124. 961	73. 557	143. 211	32.601	43.486	64. 448	43.661	36.987
				150. 983	162.248	310.059	210.563	225.764
HAMMA1002215	446.836	148. 590			35.094	50.008	34, 183	47.670
MACMA1002219	103.054	68. 338	110.047	29, 595			50.692	104. 338
MAPMA1002224	155. 329	135.036	325. 596	92.243	139, 113	54. 888		
MANYA1002229	54. 055	19. 297	24.594	8, 408	18.280	19.024	14.880	18. 482
HALMA1002230	131.172	96.706	345. 936	76.632	50. 164	62. 315	35. 205	65.871
HAMMA1002233	40. 299	20. 503	27.780	14. 545	13.380	24. 157	18.866	16.294
MANMA1002234	16.951	13.815,	19.460	7, 251	4, 128	10.631	13.812	19.438
MANMA1002236	50.642	23. 553	50. 583	14, 162	51.817	24.897	29.324	44.837
HAMMA1002243	88.955	30.943	38. 127	26.451	21.889	37, 268	32.369	10.849
		23. 851	171.031	56.513	74, 300	48. 863	11.431	66.114
MANMA 1002250	101.569			80.630	175.660	370.878	217, 429	157, 156
MAMMA1002253	515. 165	161.871	122.750			98. 387	72.076	331.998
HARMA1002267	129. 167	239.800	180.046	95.357	56.654			13.929
MAMMA 1002268	36.456	16.771	39.216	17.501	24. 043	16.873	20.704	
MANMA1002269	27.848	6.625	13.419	16.093	10.154	9.666	6.915	4.635
MALMA1002282	53.648	58. 269	178. 298	38, 160	60.059	34.106	22.977	37.892
MAMMA 1002292	62.491	17.873	48. 526	22.803	16.647	14.012	30.027	30.270
MANMA1002293	236. 280	162.513	481.000	154. 526	85. 449	104.060	60.152	54.729
MANMA 1002294	110.705	24.664	124.002	36.492	33.138	43, 853	25, 143	19.816
MALMA 1002297	66. 424	40.774	88. 229	32.940	16. 126	21.061	14. 524	17.505
MARSIA 1002298	104. 368	30.772	64, 493	24.071	29.853	40. 308	35.653	29.912
				29.656	30. 944	33.813	19,722	23. 248
MANA 1002299	102.764	41.185	67. 139		29.756	27.771	17. 935	16.223
MANMA 1002308	69. 299	30.798	86.503	30.668			183. 571	203, 149
MANSMA 1002310	494. 257	272.509	645.571	186.568	219.463	344.867		
MANSA 1002311	151.653	60.941	315.707	69.190	66.700	63.609	50.563	40.723
MARSMA 1002312	79.548	36.483	113.839	34.110	19, 878	36.852	19.114	16. 993
MAGMA 1002317	96.094	32.026	188.632	45. 170	46.365	46. 409	41.391	20. 920
MANNA 1002319	141.320	69.599	218.472	74.218	50.463	59.927	44. 261	42.418
MANMA 1002322	144. 393	65, 401	253.730	67.857	46.931	25. 375	51.002	44. 82
MANMA 1002329	49.002	17, 163	28. 349	17.067	21.239	27.218	20. 223	13.61
MALMA1002332	55. 840	30.915	137.766	47. 492	35.312	32, 956	23, 130	16.41
		17.882	32.309	19. 280	28.576	31, 145	41.629	17.63
MARMA 1002333	75. 478			54,778	40.367	18.695	38.972	26.410
MAIMA 1002335	171.866	50.373	149. 587			48.035	33.591	31.79
MALSAN 1002339	91.741	62.618	152.049	63.915	53.097	33.327	45. 235	27.50
MARMA 1002347	98.915	55, 800	120.784	40.650	55.929			19.63
HALSMA1002351	70.045	22.016	35.600	18, 133	20.122	33.583	21.722	
MANMA 1 002352	52. 143	17.786	22.690	23.069	12.412	24.411	13.818	11.94
HAC#14 1002353	128.336	52.785	144,030	46.481	46.561	36.806	12.132	34.57
MAYMA1002355	46.995	34, 505	123.684	29.737	22.025	29. 352	6.766	22.66
MANMA 1002356	40.901	21.732	86. 932	22, 189	25. 451			18. 95
		92. 529		168. 428	142.084	59.794	89.656	51.18
MANSAA 1009950						9, 982	9, 481	12.12
MANMA 1002359	276.825		A7 182	1 10 001	1 18.409			
PAPPA 1002360	42.725	25.740	47. 382	16.661	18.409			27. 24
MARSA 1002360	42.725 152.118	25.740 88.131	201.317	50.907	41.767	51.778	26.886	
HAPPA 1002360 HAPPA 1002361 HAPPA 1002362	42.725 152.118 39.281	25.740 88.131 22.692	201.317 119.094	50. 907 21. 154	41.767 14.517	51.778 23.579	26.886 14.318	19.59
PAPAA 1 002 360 MARAA 1 002 361 PAPAA 1 002 362 PAPAA 1 002 367	42.725 152.118 39.281 142.262	25.740 88.131 22.692 75.867	201.317 119.094 50.909	50. 907 21. 154 48. 285	41.767 14.517 31.065	51.778 23.579 65.479	26.886 14.318 60.201	19.59
HAPPA 1002360 HAPPA 1002361 HAPPA 1002362 HAPPA 1002367 HAPPA 1002371	42. 725 152. 118 39. 281 142. 262 119. 755	25.740 88.131 22.692 75.867 66.644	201.317 119.094 50.909 278.090	50. 907 21. 154 48. 285 138. 658	41.767 14.517 31.065 42.317	51.778 23.579 65.479 49.599	26.886 14.318 60.201 32.494	19.59 210.78 49.29
PAPAA 1 002 360 MARAA 1 002 361 PAPAA 1 002 362 PAPAA 1 002 367	42.725 152.118 39.281 142.262	25.740 88.131 22.692 75.867	201.317 119.094 50.909	50. 907 21. 154 48. 285 138. 658 38. 559	41.767 14.517 31.065 42.317 31.139	51.778 23.579 65.479 49.599 36.350	26.886 14.318 60.201 32.494 34.696	19.59 210.78 49.29 25.22
HAPPA 1002360 HAPPA 1002361 HAPPA 1002362 HAPPA 1002367 HAPPA 1002371	42.725 152.118 39.281 142.262 119.755 90.587	25.740 88.131 22.692 75.867 66.644	201.317 119.094 50.909 278.090	50. 907 21. 154 48. 285 138. 658	41.767 14.517 31.065 42.317 31.139	51. 778 23. 579 65. 479 49. 599 36. 350 40. 126	26.886 14.318 60.201 32.494 34.696 29.975	19.59 210.78 49.29 25.22 44.4
HAMA 1002360 HAMA 1002361 HAMA 1002362 HAMA 1002367 HAMA 1002371 HAMA 1002380 HAMA 1002384	42.725 152.118 39.281 142.262 119.755 90.587 90.935	25.740 88.131 22.692 75.867 66.644 47.691 85.538	201.317 119.094 50.909 278.090 161.106 249.278	50. 907 21. 154 48. 285 138. 658 38. 559	41.767 14.517 31.065 42.317 31.139 46.508	51. 778 23. 579 65. 479 49. 599 36. 350 40. 126	26.886 14.318 60.201 32.494 34.696	19.59 210.78 49.29 25.22 44.41 7.60
EACHA 1002360 EACHA 1002361 EACHA 1002362 EACHA 1002367 EACHA 1002367 EACHA 1002380 EACHA 1002384 EACHA 1002385	42.725 152.118 39.281 142.262 119.755 90.587 90.935 13.712	25,740 88,131 22,692 75,867 66,644 47,691 85,538 7,306	201.317 119.094 50.909 278.090 161.106 249.278 6.051	50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420	41.767 14.517 31.065 42.317 31.139 46.508 3.720	51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699	26.886 14.318 60.201 32.494 34.696 29.975	19.59 210.78 49.29 25.22 44.41 7.60
EACHA 1002360 EACHA 1002351 EACHA 1002362 EACHA 1002367 EACHA 1002367 EACHA 1002371 EACHA 1002384 EACHA 1002385 EACHA 1002385	42.725 152.118 39.281 142.262 119.755 90.587 90.935 13.712 119.086	25.740 88.131 22.692 75.867 66.644 47.691 85.538 7.306 26.468	201.317 119.094 50.909 278.090 161.106 249.278 6.051 66.535	50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420 12. 989	41.767 14.517 31.065 42.317 31.139 46.508 3.720 40.464	51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956	26.886 14.318 60.201 32.494 34.696 29.975 8.116	19.59 210.78 49.29 25.22 44.41 7.60
EAMA 1002360 EAMA 1002351 EAMA 1002362 EAMA 1002367 EAMA 1002367 EAMA 1002381 EAMA 1002381 EAMA 1002382 EAMA 1002383 EAMA 1002383	42, 725 152, 118 39, 281 142, 262 119, 755 90, 587 90, 935 13, 712 119, 086 90, 573	25,740 88,131 22,692 75,867 66,644 47,691 85,538 7,306 26,468 32,273	201.317 119.094 50.909 278.090 161.106 249.278 6.051 66.535 97.224	50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420 12. 989 19. 547	41.767 14.517 31.065 42.317 31.139 46.508 3.720 40.464 21.438	51.778 23.579 65.479 49.599 36.350 40.126 9.699 53.956 26.503	26.886 14.318 60.201 32.494 34.696 29.975 8.116 37.080 20.868	19.59 210.78 49.25 25.22 44.41 7.60 19.51
EACHA 1002360 EACHA 1002351 EACHA 1002362 EACHA 1002367 EACHA 1002367 EACHA 1002371 EACHA 1002384 EACHA 1002385 EACHA 1002385	42.725 152.118 39.281 142.262 119.755 90.587 90.935 13.712 119.086	25.740 88.131 22.692 75.867 66.644 47.691 85.538 7.306 26.468	201.317 119.094 50.909 278.090 161.106 249.278 6.051 66.535 97.224	50. 907 21. 154 48. 285 138. 658 38. 559 71. 113 7. 420 12. 989 19. 547	41.767 14.517 31.065 42.317 31.139 46.508 3.720 40.464 21.438 82.112	51. 778 23. 579 65. 479 49. 599 36. 350 40. 126 9. 699 53. 956 26. 503 77. 745	26.886 14.318 60.201 32.494 34.696 29.975 8.116 37.080 20.868 28.921	14. 25 53. 90

Table 59

Section   Content   Cont		HAPMA1002400	10.797	7.113	11, 587	4.041	5. 847	4.732	4.516	4. 194
HARMANIO   1991   1995   1986   1987   198										
### CHAPMIND (24) 13 193, 065										
SUCY_SINDOZ417   30.916   25.195   58.136   15.593   17.649   14.266   7.765   11.985   1.685   10.2	5									
SMESTANDOZ477   87.721   47.715   208.529   48.123   38.191   40.117   26.156   31.585   31.585   37.126   38.231   32.331   39.146   38.253   34.156   37.126   38.231   38.231   39.146   38.253   34.156   37.126   38.231   38.231   39.146   38.253   34.156   37.126   38.231   39.231   39.231   39.231   39.231   36.229   25.042   39.231   39.231   39.255   34.156   37.255   35.246   39.231   39										
NASAN 1002448   17.152   72.024   272.113   68.594   58.156   51.766   57.518   57.126										
NUM_NICO   19.0   30.843   21.726   38.763   19.586   19.585   44.397   36.529   25.042     NUM_NICO   22446   102.855   36.748   90.796   22.555   36.516   45.916   45.916     NUM_NICO   2447   77.962   49.457   17.445   42.555   21.446   36.510   25.92   27.967     NUM_NICO   2447   77.962   49.457   17.445   42.555   21.446   36.510   25.92   27.967     NUM_NICO   2447   77.962   49.457   17.445   42.555   21.446   36.510   25.92   27.967     NUM_NICO   2447   77.962   49.457   17.445   42.555   21.446   36.510   25.92   27.967     NUM_NICO   2447   27.967   39.505   53.9572   188.845   118.797   99.596   72.794   101.951     NUM_NICO   2447   24.967   34.520   44.84   72.551   25.745   31.859   67.395   41.900   28.713     NUM_NICO   2448   97.969   94.520   44.464   18.727   24.4045   05.577   30.577   24.7045   05.657   27.107     NUM_NICO   2447   95.967   94.520   44.464   18.727   24.4045   05.657   27.107   17.415     NUM_NICO   2476   52.777   10.542   97.707   23.707   24.685   10.963   12.591   28.386     NUM_NICO   2476   53.982   26.009   77.707   23.707   24.685   10.963   12.591   26.386     NUM_NICO   2486   25.602   25.6135   77.461   32.376   24.865   10.963   12.591   26.386     NUM_NICO   2486   25.602   25.6135   77.461   32.376   24.865   10.963   12.591   26.386     NUM_NICO   2486   25.602   26.2135   77.461   32.376   24.985   10.963   17.311   49.943     NUM_NICO   2486   25.602   26.2135   77.461   32.376   24.985   26.986   17.311   49.943     NUM_NICO   2486   25.602   27.727   24.662   24.255   29.322   31.125   26.896   19.794   5.551     NUM_NICO   25.602   26.602   27.727   27.864   27.25   27.727   27.70										
NUMANIA   NUMA										
SEM_MAIN 1002446   102.855   36.748   90.796   22.955   36.351   49.598   42.676   12.897										
GMEMATO07445   145 00   201 950 539 572   184 845   18,797   99, 595   77,794   103, 951	10									
SHEEN 10074-65   314. 500   201. 950   539. 572   188. 845   118. 797   99. 956   72. 794   101. 951     HARPEN 10074-65   204. 658   47. 899   13. 552   28. 137   55. 948   63. 968   52. 524   64. 401     HARPEN 10074-65   204. 659   47. 899   48. 48. 48   48. 573   24. 64. 55   50. 857   37. 103   17. 415     HARPEN 10072-65   34. 579   34. 529   44. 48. 48   48. 573   24. 64. 55   50. 857   37. 103   17. 415     HARPEN 10072-67   56. 277   10. 542   19. 523   14. 778   99. 364   20. 022   17. 241   15. 392     HARPEN 10072-77   55. 277   10. 542   19. 523   14. 778   99. 364   20. 022   17. 241   15. 392     HARPEN 10072-88   25. 50. 98   77. 707   23. 670   24. 685   10. 653   12. 591   26. 386     HARPEN 10072-88   25. 50. 74   56. 245   77. 707   23. 670   24. 685   10. 653   12. 591   26. 386     HARPEN 10072-88   25. 50. 27   24. 285   12. 392   31. 125   20. 038   77. 311   49. 343     HARPEN 10072-88   25. 30. 27   23. 81   164. 418   25. 376   48. 947   43. 135   11. 733   44. 491     HARPEN 10072-89   36. 729   79. 24. 265   79. 292   31. 125   25. 356   79. 74   44. 45     HARPEN 10072-89   36. 729   79. 293   79. 720   79. 724   74. 135   77. 311   49. 343     HARPEN 10072-89   36. 729   79. 293   79. 720   79. 724   74. 135   77. 731   49. 343     HARPEN 10072-89   36. 727   79. 724   79. 727   79. 727   79. 727   79. 728   79.										
Hammodia   1946   1947   1948   1947   1948   194		MAMMA 1002441								
BMEMANDOZASS   130. 489										
BACCAN 1002466										
Section   Color   Co										
SHEPSIN 1002470   \$6, 277   10, 542   19, 523   14, 778   5, 384   20, 072   21, 241   15, 324	15									
BLASSA 1007475   35.982   26.009   77.707   27.670   24.885   10.963   12.591   26.386   BLASSA 1007485   255.024   45.191   44.99   40.755   50.788   48.101   35.187   30.088   BLASSA 1007485   255.024   56.749   23.381   164.418   25.376   48.947   43.381   17.331   49.943   14.2491   1007484   55.072   20.346   24.725   72.935   13.125   28.980   19.794   5.551   14.2481   1007524   73.678   20.346   24.725   72.932   13.125   28.980   19.794   5.551   14.2481   1007524   73.578   20.346   24.725   72.932   13.125   28.980   19.794   5.551   14.2481   1007524   73.14   15.000   70.000   32.939   72.173   54.25   25.458   27.749   12.355   14.645										
HAMMAT NODZAGE										
MARMAN 10072486   555. 074   55. 215   75. 461   37. 978   72. 095   20. 038   77. 311   49. 943										
Hamman 1007249										
### PAMPAN 1002489										
HAMMATIOUZSSI   71, 628   20, 842   11, 923   21, 047   20, 268   27, 749   12, 366   14, 645	20									
### PAREMATORYSTO 82,789 19.903 43,603 113,551 9,151 28,515 27,989 12,505 192600251 101,182 27,725 28,460 21,181 31,900 45,227 63,30 25,658 120,2251 131,415 100,020 322,993 72,173 54,265 23,145 30,820 51,328 120,2251 131,415 100,020 322,993 72,173 54,265 23,145 30,820 51,328 120,2251 131,415 100,020 322,993 72,173 54,265 23,145 30,820 51,328 120,2251 131,415 120,0256 131,527 31,033 30,923 62,549 16,548 18,644 38,344 32,052 17,411 132,225 120,	~~									
MARSATOD2538   101.   162   27.   725   78.   460   21.   161   31.   900   45.   529   26.   380   25.   558   104881002545   31.   415   100.020   322.   953   72.   173   54.   255   23.   145   30.   820   51.   328   104881002556   51.   318   30.   820   51.   328   32.   325   32.   325   32.   325   32.   325   32.   325   32.   325   32.   325   32.   33.   32.   32.   32.   32.   33.   32.   32.   32.   33.   32.   32.   33.   32.   32.   33.   32.   33.   32.   33.   32.   33.   32.   33.   32.   33.   32.   33.   3										
MARSA   1002545   131. 415   100. 020   322. 993   72. 173   54. 265   23. 145   30. 820   51. 328										
NAMESA 1002556   51,033   30,932   62,549   16,548   18,644   38,344   32,052   17,411										
HAMMATODZSGS   199.748   128.004   556.958   135.854   118.280   54.740   81.217   51.556	25									
HAMMAT 1002565	23									
HAMMAID02566   29.155   16.405   7.906   3.460   1.967   13.518   5.709   5.318										
HAMMA1002571   73.034   22.187   37.154   25.594   6.079   28.030   19.946   20.955     HAMMA1002573   218.479   62.569   183.544   61.350   46.029   113.781   65.617   60.521     HAMMA1002584   199.621   18.498   33.802   10.617   22.615   43.283   55.199   26.452     HAMMA1002584   244.467   197.526   384.879   79.185   103.251   112.917   113.914   151.642     HAMMA1002585   133.865   28.963   56.983   17.185   103.251   112.917   113.914   151.642     HAMMA1002585   67.168   39.043   44.767   15.656   19.252   29.556   35.555   19.945     HAMMA1002589   98.170   25.567   26.638   16.923   18.956   18.249   16.364   12.591     HAMMA1002589   268.176   57.804   202.329   36.276   77.487   180.923   123.883   42.552     HAMMA1002587   76.091   50.352   131.097   33.605   42.551   25.425   36.396   34.764     HAMMA1002587   76.091   50.352   131.097   33.605   42.551   25.425   36.396   34.764     HAMMA1002588   69.190   45.133   59.324   58.225   35.339   68.531   47.164   70.246     HAMMA1002587   36.319   11.631   557.754   135.485   46.672   98.075   64.732   66.103     HAMMA1002618   30.423   66.208   129.807   53.454   46.996   53.758   43.899   55.854     HAMMA1002619   34.076   14.223   23.292   10.350   14.540   15.236   11.009   118.254     HAMMA1002625   112.756   60.308   129.807   53.454   46.996   53.758   43.899   55.854     HAMMA1002627   12.756   60.308   129.807   53.454   46.996   53.758   43.899   55.854     HAMMA1002627   12.756   60.308   129.807   53.454   46.996   53.758   43.899   55.854     HAMMA1002633   50.470   10.960   11.524   64.401   102.216   102.611   54.682   73.325     HAMMA1002633   50.470   10.960   11.524   65.679   37.741   10.19   10.741   11.301     HAMMA1002635   59.898   50.529   142.123   25.014   15.348   18.150   38.497   28.162     HAMMA1002635   59.898   50.529   142.123   25.014   15.348   18.150   38.497   28.162     HAMMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.497   28.162     HAMMA1002636   59.898   50.529   142.123   25.014   1										
Chaman   Colorest										
CLABMATOUZESTS   109, 621   18, 498   33, 802   10, 617   22, 615   43, 283   55, 199   26, 452								113, 781	65.617	
ENMANDESIMA   244, 467   197, 626   384, 879   79, 185   103, 251   112, 917   113, 914   151, 642	20							43. 283	55. 199	26.452
CAMMAN   COURSE   133   865   28   963   56   983   17   186   16   306   13   727   51   687   25   753	30								113.914	151.642
CAMMA 1002586   67.168   39.043   34.776   15.656   19.252   29.596   35.555   19.945									51.687	
HAMMA1002589   98. 120   25. 567   26. 638   16. 923   18. 956   18. 249   16. 364   12. 591		MANNA 1002586						29. 596	35. 555	
HABBIA1002590   268. 176   57. 804   202. 329   36. 276   77. 487   180. 923   123. 883   42. 552     HABBIA1002593   131. 425   64. 951   130. 257   54. 131   23. 515   55. 983   37. 410   36. 272     HABBIA1002598   69. 190   45. 133   59. 324   58. 225   35. 339   68. 531   47. 164   70. 246     HABBIA1002603   122. 932   40. 124   155. 801   51. 386   48. 672   98. 075   64. 732   66. 103     HABBIA1002617   363. 139   211. 631   557. 754   145. 485   146. 260   203. 052   110. 009   118. 254     HABBIA1002618   90. 423   66. 208   129. 807   53. 454   46. 096   53. 758   43. 899   55. 854     HABBIA1002619   34. 076   14. 223   23. 292   10. 350   14. 540   15. 236   12. 465   13. 642     HABBIA1002623   89. 689   68. 083   149. 811   64. 401   102. 216   102. 611   54. 682   73. 325     HABBIA1002623   89. 689   68. 083   149. 811   64. 401   102. 216   102. 611   54. 682   73. 325     HABBIA1002627   9. 090   2. 616   7. 631   2. 675   0. 000   3. 940   7. 852   8. 826     HABBIA1002631   50. 470   10. 960   11. 524   6. 679   3. 741   10. 219   10. 741   11. 301     HABBIA1002633   59. 898   50. 529   142. 123   25. 014   15. 348   18. 150   38. 018   22. 608     HABBIA1002635   59. 898   50. 529   142. 123   25. 014   15. 348   18. 150   38. 018   22. 608     HABBIA1002655   49. 661   48. 800   69. 217   43. 621   54. 730   39. 438   38. 742   48. 014     HABBIA1002655   49. 661   48. 800   69. 217   43. 621   54. 730   39. 438   38. 742   48. 014     HABBIA1002655   49. 661   48. 800   69. 217   43. 621   54. 730   39. 438   38. 742   48. 014     HABBIA1002656   212. 410   44. 430   94. 935   34. 850   32. 770   58. 417   41. 476   33. 910     HABBIA1002655   49. 661   48. 800   69. 217   43. 621   54. 730   39. 438   38. 742   48. 014     HABBIA1002655   49. 661   48. 800   69. 217   43. 621   54. 730   39. 438   38. 742   48. 014     HABBIA1002656   212. 410   44. 430   94. 935   34. 850   32. 770   58. 417   41. 476   33. 910     HABBIA1002665   23. 6. 733   190. 056   600. 904   183. 784   112.								18. 249	16. 364	12.591
HAMMA 1002593   131, 425   64, 951   130, 257   54, 131   23, 515   55, 983   37, 410   36, 272     HAMMA 1002598   69, 190   45, 133   59, 324   58, 225   35, 339   68, 531   47, 164   70, 246     HAMMA 1002603   122, 932   40, 124   155, 801   51, 386   48, 672   98, 075   64, 732   66, 103     HAMMA 1002612   330, 999   152, 583   441, 574   105, 603   112, 764   175, 106   98, 853   99, 475     HAMMA 1002613   363, 139   211, 631   557, 754   145, 485   146, 260   203, 052   110, 009   118, 254     HAMMA 1002619   34, 076   14, 223   23, 292   10, 350   14, 540   15, 236   12, 465   13, 642     HAMMA 1002629   34, 076   14, 223   23, 292   10, 350   14, 540   15, 236   12, 465   13, 642     HAMMA 1002622   112, 756   60, 308   263, 518   46, 461   43, 508   41, 984   32, 044   52, 630     HAMMA 1002623   89, 689   68, 083   149, 811   64, 401   102, 216   102, 611   54, 682   73, 325     HAMMA 1002627   9, 090   2, 616   7, 631   2, 675   0, 000   3, 940   7, 852   8, 826     HAMMA 1002627   9, 090   2, 616   7, 631   2, 675   0, 000   3, 940   7, 852   8, 826     HAMMA 1002623   50, 470   10, 960   11, 524   6, 753   3, 741   10, 219   10, 741   11, 301     MAMMA 1002633   32, 234   20, 386   37, 729   16, 053   9, 358   12, 456   8, 681   32, 169     HAMMA 1002636   59, 898   50, 529   142, 123   25, 014   15, 348   18, 150   38, 018   22, 608     HAMMA 1002637   58, 583   21, 541   11, 323   5, 892   14, 789   18, 069   26, 406   27, 104     HAMMA 1002636   59, 898   50, 529   142, 123   25, 014   15, 348   18, 150   38, 018   22, 608     HAMMA 1002637   58, 583   21, 541   11, 323   5, 892   14, 789   18, 069   26, 406   27, 104     HAMMA 1002638   59, 898   50, 529   142, 123   25, 014   15, 348   18, 150   38, 018   22, 608     HAMMA 1002638   59, 698   69, 556   44, 999   60, 882   59, 089   42, 135   62, 414   54, 651     HAMMA 1002662   122, 410   44, 430   94, 935   34, 850   32, 770   58, 417   41, 476   39, 910     HAMMA 1002665   236, 733   190, 056   600, 904   183, 784   112, 684   133, 133			268. 176	57.804		36.276		180. 923	123.883	42.552
HAMMA1002598   Feb.   Feb.   Feb.   Feb.   Feb.   HAMMA1002598   Feb.							23.515	55. 983	37.410	36.272
HAMMA1002603 122.932 40.124 155.801 51.386 48.672 98.075 64.732 66.103 HAMMA1002612 330.999 152.583 441.574 105.603 112.764 175.106 98.853 99.475 HAMMA1002617 363.139 211.631 557.754 145.485 146.260 203.052 110.009 118.254 HAMMA1002618 90.423 66.208 129.807 53.454 46.096 53.758 43.899 55.854 HAMMA1002622 112.756 60.308 263.518 46.461 43.508 41.984 32.044 52.630 HAMMA1002623 89.689 68.083 149.811 64.401 102.216 102.611 54.682 73.325 HAMMA1002623 89.689 68.083 149.811 64.401 102.216 102.611 54.682 73.325 HAMMA1002627 9.090 2.616 7.631 2.675 0.000 3.940 7.852 8.826 HAMMA1002627 9.090 2.616 7.631 2.675 0.000 3.940 7.852 8.826 HAMMA1002631 50.470 10.960 11.524 6.679 3.741 10.219 10.741 11.301 MAMMA1002635 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002637 58.583 21.541 11.323 5.882 14.789 18.0659 26.406 22.104 HAMMA1002638 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HAMMA1002658 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HAMMA1002658 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HAMMA1002658 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HAMMA1002658 21.2.410 44.430 94.935 34.850 32.770 58.417 41.476 39.910 HAMMA1002655 236.733 190.056 600.904 183.784 112.684 133.133 101.570 153.389 HAMMA1002657 89.496 41.623 60.274 25.563 20.577 26.452 50.459 40.518	35	MANMA1002597	76.091	50. 352	131.097		42.551		36. 396	34.764
HAPMA1002612   330,999   152,583   441,574   105,603   112,764   175,106   98,853   99,475     HAPMA1002617   363,139   211,631   557,754   145,485   146,260   203,052   110,009   118,254     HAPMA1002618   90,423   66,208   129,807   53,454   46,096   53,758   43,899   55,854     HAPMA1002619   34,076   14,223   23,292   10,350   14,540   15,236   12,465   13,642     HAPMA1002622   112,755   60,308   263,518   46,461   43,508   41,984   32,044   52,630     HAPMA1002623   89,689   68,083   149,811   64,401   102,216   102,611   54,682   73,325     HAPMA1002625   83,660   44,949   94,038   26,154   32,540   34,576   38,497   28,162     HAPMA1002627   9,090   2,616   7,631   2,675   0,000   3,940   7,852   8,826     HAPMA1002631   50,470   10,960   11,524   6,679   3,741   10,219   10,741   11,301     HAPMA1002633   59,898   50,529   142,123   25,014   15,348   18,150   38,018   22,608     HAPMA1002637   58,583   21,541   11,323   5,892   14,789   18,069   26,406   22,104     HAPMA1002638   59,898   50,529   142,123   25,014   15,348   18,150   38,018   22,608     HAPMA1002638   55,442   29,770   36,308   23,176   15,750   18,816   26,997   38,809     HAPMA1002650   15,384   6,907   9,595   4,820   3,958   6,140   8,225   6,042     HAPMA1002655   49,661   48,800   69,217   43,621   64,730   39,438   38,742   48,014     HAPMA1002655   49,667   25,105   13,568   1,569   8,462   23,347   10,991   22,157     HAPMA1002655   49,617   25,105   13,568   1,569   8,462   23,347   10,991   22,157     HAPMA1002655   236,733   190,056   600,904   183,784   112,684   133,133   101,570   153,389     HAPMA1002665   236,733   190,056   600,904   183,784   112,684   133,133   101,570   153,389     HAPMA1002671   89,496   41,623   60,274   25,563   20,577   26,452   50,459   40,518				45. 133				68.531		
HAMMA1002817 363.139 211.631 557.754 145.485 146.260 203.052 110.009 118.254 HAMMA1002618 90.423 66.208 129.807 53.454 46.096 53.758 43.899 55.854 HAMMA1002619 34.076 14.223 23.292 10.350 14.540 15.236 12.465 13.642 HAMMA1002623 89.689 68.083 149.811 64.401 102.216 102.611 54.682 73.325 HAMMA1002625 83.660 44.949 94.038 26.154 32.540 34.576 38.497 28.162 HAMMA1002627 9.090 2.616 7.631 2.675 0.000 3.940 7.852 8.826 HAMMA1002627 9.090 2.616 7.631 2.675 0.000 3.940 7.852 8.826 HAMMA1002629 111.050 96.279 397.433 77.573 45.933 89.752 53.737 108.399 10.8240 10.960 11.524 6.679 3.741 10.219 10.741 11.301 HAMMA1002631 50.470 10.960 11.524 6.679 3.741 10.219 10.741 11.301 HAMMA1002633 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002637 58.583 21.541 11.323 5.892 14.789 18.069 26.406 22.104 HAMMA1002634 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HAMMA1002655 15.384 6.907 9.595 44.820 3.958 6.140 8.225 6.042 10.44 11.301 11.30					155.801					
HAMMA1002618 90.423 66.208 129.807 53.454 46.096 53.758 43.899 55.854 HAMMA1002619 34.076 14.223 23.292 10.350 14.540 15.236 12.465 13.642 HAMMA1002622 112.756 60.308 263.518 46.461 43.508 41.984 32.044 52.630 HAMMA1002623 89.689 68.083 149.811 64.401 102.216 102.611 54.682 73.325 HAMMA1002625 83.660 44.949 94.038 26.154 32.540 34.576 38.497 28.162 HAMMA1002627 9.090 2.616 7.631 2.675 0.000 3.940 7.852 8.826 HAMMA1002629 111.050 96.279 397.433 77.573 45.933 89.752 53.737 108.399 HAMMA1002631 50.470 10.960 11.524 6.679 3.741 10.219 10.741 11.301 MAMMA1002633 32.234 20.386 37.729 16.053 9.358 12.456 8.681 32.169 HAMMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HAMMA1002636 49.661 29.770 36.308 23.176 15.750 18.816 26.997 38.809 HAMMA1002658 49.661 29.770 36.308 23.176 15.750 18.816 26.997 38.809 HAMMA1002658 49.661 29.770 36.308 23.176 15.750 18.816 26.997 38.809 HAMMA1002658 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HAMMA1002655 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HAMMA1002665 236.733 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HAMMA1002667 236.733 190.058 600.904 183.784 112.684 133.133 101.570 153.389			330. 999			105.603	112.764			
HANMATOUZESS 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HANMATOUZESS 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HANMATOUZESS 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HANMATOUZESS 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HANMATOUZESS 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HANMATOUZESS 53.63 19.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 19.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.63 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.673 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.673 190.058 600.904 183.784 112.684 133.133 101.570 153.389 HANMATOUZESS 23.673 190.058 600.904 183.784 112.684 133.133 101.570 153.389										
HARMA1002622   112.756   50.308   263.518   46.461   43.508   41.984   32.044   52.630     HARMA1002623   89.689   68.083   149.811   64.401   102.216   102.611   54.682   73.325     HARMA1002625   83.660   44.949   94.038   26.154   32.540   34.576   38.497   28.162     HARMA1002627   9.090   2.616   7.631   2.675   0.000   3.940   7.852   8.826     HARMA1002629   111.050   96.279   397.433   77.573   45.933   89.752   53.737   108.399     HARMA1002631   50.470   10.960   11.524   6.679   3.741   10.219   10.741   11.301     HARMA1002633   32.234   20.386   37.729   16.053   9.358   12.456   8.681   32.169     HARMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.018   22.608     HARMA1002637   58.583   21.541   11.323   5.892   14.789   18.069   26.406   22.104     HARMA1002638   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     HARMA1002638   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HARMA1002654   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HARMA1002655   61.935   69.556   44.994   60.882   59.089   42.135   62.414   54.651     HARMA1002655   236.733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HARMA1002667   22.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HARMA1002667   22.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HARMA1002667   22.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HARMA1002667   22.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HARMA1002667   89.495   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002667   23.6733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HARMA1002667   89.495   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002667   23.6733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HARMA1002667   236.773   236.774   25.563   20.577   26.452   50.4	40									
HAPMA1002623	40									
HARMA1002627   9.090   2.616   7.631   2.675   0.000   3.940   7.852   8.826     HARMA1002627   9.090   2.616   7.631   2.675   0.000   3.940   7.852   8.826     HARMA1002629   111.050   96.279   397.433   77.573   45.933   89.752   53.737   108.399     HARMA1002631   50.470   10.960   11.524   6.679   3.741   10.219   10.741   11.301     MARMA1002633   32.234   20.386   37.729   16.053   9.358   12.456   8.681   32.169     HARMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.018   22.608     HARMA1002637   58.883   21.541   11.323   5.892   14.789   18.069   26.406   22.104     HARMA1002648   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     HARMA1002648   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HARMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     HARMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     HARMA1002662   122.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HARMA1002665   236.733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HARMA10										
HAMMA1002627   9.090   2.616   7.631   2.675   0.000   3.940   7.852   8.826     HAMMA1002629   111.050   96.279   397.433   77.573   45.933   89.752   53.737   108.399     HAMMA1002631   50.470   10.960   11.524   6.679   3.741   10.219   10.741   11.301     HAMMA1002633   32.234   20.386   37.729   16.053   9.358   12.456   8.681   32.169     HAMMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.018   22.608     HAMMA1002637   58.583   21.541   11.323   5.892   14.789   18.069   26.406   22.104     HAMMA1002637   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     HAMMA1002636   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HAMMA1002655   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HAMMA1002655   61.935   69.556   64.994   60.882   59.089   42.135   62.414   54.651     HAMMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     HAMMA1002662   122.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HAMMA1002665   236.733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     HAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518										
HARMA1002639 111.050 96.279 397.433 77.573 45.933 89.752 53.737 108.399 HARMA1002631 50.470 10.960 11.524 6.679 3.741 10.219 10.741 11.301 HARMA1002633 32.234 20.386 37.729 16.053 9.358 12.456 8.681 32.169 HARMA1002636 59.898 50.529 142.123 25.014 15.348 18.150 38.018 22.608 HARMA1002637 58.583 21.541 11.323 5.892 14.789 18.069 26.406 22.104 HARMA1002645 55.442 29.770 36.308 23.176 15.750 18.816 26.997 38.809 HARMA1002656 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014 HARMA1002655 15.384 5.907 9.595 4.820 3.958 6.140 8.225 6.042 HARMA1002655 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HARMA1002655 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HARMA1002662 122.410 44.430 94.935 34.850 32.770 58.417 41.476 39.910 HARMA1002665 236.733 190.056 600.904 183.784 112.684 133.133 101.570 153.389										
HAPMA1002631   50.470   10.960   11.524   6.679   3.741   10.219   10.741   11.301     MARMA1002633   32.234   20.385   37.729   16.053   9.358   12.456   8.681   32.169     MARMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.018   22.608     MARMA1002637   58.583   21.541   11.323   5.892   14.789   18.069   26.406   22.104     MARMA1002645   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     MARMA1002655   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     MARMA1002650   15.384   6.907   9.595   4.820   3.958   6.140   8.225   6.042     MARMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     MARMA1002662   122.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     MARMA1002665   236.733   190.058   600.904   183.784   112.684   133.133   101.570   153.389     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518     MARMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518		The state of the s	<del></del>							
MAMMA1002633   32,234   20,386   37,729   16,053   9,358   12,456   8,681   32,169										
HAMMA1002636   59.898   50.529   142.123   25.014   15.348   18.150   38.018   22.608     HAMMA1002637   58.583   21.541   11.323   5.892   14.789   18.069   26.406   22.104     HAMMA1002646   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     HAMMA1002648   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     HAMMA1002650   15.384   6.907   9.595   4.820   3.958   6.140   8.225   6.042     HAMMA1002652   61.935   69.556   44.994   60.882   59.089   42.135   62.414   54.651     HAMMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     HAMMA1002662   122.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     HAMMA1002665   236.733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     HAMMA1002671   89.495   41.623   60.274   25.563   20.577   26.452   50.459   40.518	45									
MAMMA1002637   58.583   21.541   11.323   5.892   14.789   18.069   26.406   22.104     MAMMA1002645   55.442   29.770   36.308   23.176   15.750   18.816   26.997   38.809     MAMMA1002648   49.661   48.800   69.217   43.621   64.730   39.438   38.742   48.014     MAMMA1002650   15.384   6.907   9.595   4.820   3.958   6.140   8.225   6.042     MAMMA1002652   61.935   69.556   44.994   60.882   59.089   42.135   62.414   54.651     MAMMA1002655   49.617   25.105   13.568   11.569   8.462   23.347   10.991   22.157     MAMMA1002662   122.410   44.430   94.935   34.850   32.770   58.417   41.476   39.910     MAMMA1002665   236.733   190.056   600.904   183.784   112.684   133.133   101.570   153.389     MAMMA1002671   89.496   41.623   60.274   25.563   20.577   26.452   50.459   40.518										
HAMMA1002648 55.442 29.770 36.308 23.176 15.750 18.816 26.997 38.809 HAMMA1002648 49.661 48.800 69.217 43.621 64.730 39.438 38.742 48.014  HAMMA1002650 15.384 6.907 9.595 4.820 3.958 6.140 8.225 6.042 HAMMA1002652 61.935 69.556 44.994 60.882 59.089 42.135 62.414 54.651 HAMMA1002655 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HAMMA1002662 122.410 44.430 94.935 34.850 32.770 58.417 41.476 39.910 HAMMA1002665 236.733 190.056 600.904 183.784 112.684 133.133 101.570 153.389 HAMMA1002671 89.496 41.623 60.274 25.563 20.577 26.452 50.459 40.518										
MAMMA1002648		CAE 41002637								
15.384										
HAMMA1002652 61.935 69.556 44.994 60.882 59.089 42.135 62.414 54.651 HAMMA1002655 49.617 25.105 13.568 11.569 8.462 23.347 10.991 22.157 HAMMA1002662 122.410 44.430 94.935 34.850 32.770 58.417 41.476 39.910 HAMMA1002665 236.733 190.056 600.904 183.784 112.684 133.133 101.570 153.389 HAMMA1002671 89.496 41.623 60.274 25.563 20.577 26.452 50.459 40.518										
HAMMA1002655       49.617       25.105       13.568       11.569       8.462       23.347       10.991       22.157         HAMMA1002662       122.410       44.430       94.935       34.850       32.770       58.417       41.476       39.910         HAMMA1002665       236.733       190.058       600.904       183.784       112.684       133.133       101.570       153.389         HAMMA1002671       89.496       41.623       60.274       25.563       20.577       26.452       50.459       40.518	50									
HAPMA1002662       122.410       44.430       94.935       34.850       32.770       58.417       41.476       39.910         HAPMA1002665       236.733       190.056       600.904       183.784       112.684       133.133       101.570       153.389         NAPMA1002671       89.496       41.623       60.274       25.563       20.577       26.452       50.459       40.518										
HAPMA1002665 236.733 190.056 600.904 183.784 112.684 133.133 101.570 153.389  MAPMA1002671 89.496 41.623 60.274 25.563 20.577 26.452 50.459 40.518										
MAMMA1002671 89.496 41.623 60.274 25.563 20.577 26.452 50.459 40.518										
55		LIAT 14 1002671	89.496	41.623	bu. 2/4	25.563	20.5//	20.452	j 30. 459	1 40. 318
	55									

## Table 60

	MAMMA 1002673	94. 294	135.347	302.435	85.978	116.544	122.876	58.765	72.402
	HALMA1002684	169.486	32.550	60.424	32.013	39. 987	86. 564	80, 699	45.058
	MAMMA1002685	25. 020	18. 401	21.785	11.312	11.628	3. 402	5.660	25.002
	MAPMA1002692	7.274	9.361	3, 697	10.386	2.003	4, 100	3. 302	9.849
	HAPMA1002693		52. 339	15.641	32.934	10.671	20.167	32.429	30.795
		66.711							
	MAZMA1002698	39. 272	32.200	43.657	33.153	4. 354	11.796	12.328	34. 409
	MAPMA1002699	18, 348	10.645	5. 272	3. 333	2.314	3, 625	12.679	6.883
							57, 919		
	MAPMA1002701	66, 193	107. 821	326.150	82.189	33.993		29.820	56.144
	HAMMA1002708	232.250	119.730	163.846	75.850	65. 245	75.116	103.624	109.697
	MAYSMA1002711	128.862	101.834		105. 535	79. 020	76.543	26. 135	61. 975
	MAYMA1002712	55. 151	50.304	36.811	8.507	18. 857	25.978	44. 085	47.001
	MAMMA1002716	32.821	37, 741	37.674	23.554	13, 366	39, 383	49.740	33.088
				360.515	86.920	49.826	57.925	48, 421	76.576
	MAMMA 1002721	128.620	78.060						
	MAMMA 1002723	67.425	45.775	59.116	53.954	27.853	31.646	28.039	37.993
	MAPMA1002727	4.194	5.317	4.081	4.586	3.879	1.679	6.885	6. 203
				134, 784	49.359	17. 238	32.733	26. 228	67.828
	MAMMA1002728	45. 508	63.239						
	MAPMA1002742	486.871	191.088	183.567	79.031	108.740	257. 374	156.771	126. 280
	MAMMA1002743	17, 914	25.779	65.317	19.354	14.843	12. 214	24. 184	22. 277
						40.023	33, 273	23. 675	53, 991
	MAMMA1002744	70.172	65. 1841	190.550	59.599				
	MAKEMA1002746	14.967	8. 271	6. 293	9.116_	3. 957	9, 800	1.039	7.011
	MAMMA 1002748	53, 355	180.966	171.425	25.271	3.510	13.742	11.775	23.747
	MALMA 1002754	64. 093	69, 489	189. 499	44.022	29. 371	15,039	15.857	30, 299
									9. 173
•	MANMA 1002758	25.835	7.240	9.756	5. 507	5. 640	9, 500	11.968	
	MAPMA1002762	65.824	58. 122	104. 988	33.940	18.698	86.679	92.471	84.012
	MAMMA1002764	104, 828	95.058	295.803	59.465	52.006	47, 508	45.629	48. 337
								22. 519	36.212
	MARMA1002765	81.926	54. 425	185.685	56.838	25. 634	30.254		
	MANMA 1002769	20.078	9.062	33.997	9.878	15. 366	12.293	19.431	15.797
	MALMA1002771	92.652	248.038		106. 297	36. 324	95. 235	52.022	929.910
	MAURIA 1002775	51.236	37.084	125. 540	30.088	37. 975	21.242	25.695	24. 387
	MANMA 1002780	23.190	24.572	73.778	29.564	12.337	13, 199	6.027	19. 175
	MANA 1002782	76.728	28.066	76.753	28.366	26. 053	26.045	13.885	33.944
	MANIBA 1002795	17.412	3. 178	14.907	9.264	2.359	6.615	10.186	19.921
	MANMA 1002796	28. 596	28. 390	48.340	13.930	16.360	14. 274	13.494	19.709
	MARMA1002805	25. 198	16.430	30. 126	13.856	9.933	47.769	23.312	13.432
									30.489
	MAMMA1002806	84, 431	28. 564	34. 957	32.528	49. 335	29.125	31.705	
	MARMA 1002807	64, 374	42. 471	124.060	39.454	51.288	34.538	23. 265	46. 125
	MANA 1002814	28.078	31.573	133.666	35.466	14.707	19.459	22.590	33. 539
								8.038	11, 12B
	MALMA 1002817	8.719	10. 443	6.527	4.036	1.155	2.240		
	MANMA 1002820	15, 173	5.049	24.747	14.605	7.416	9.432	16.038	5, 111
	MANMA 1002830	91.438	212.662	185.761	75.492	49. 491	111.835	311.632	133. 132
			1 212.002					25.094	46.080
	MANMA1002833	90.875	71.138	237. 238	50.346	44.689	47.222		
	MAMMA1002835	28, 488	23.244	28. 102	14. 935	9. 604	12.597	16.302	12.709
	MANMA1002838	84.752	56.692	166. 200	49.694	30. 237	32.930	11.628	26.416
					23.902	32.033	41.236	27.950	47.227
	MANNA1002842	98.706	53.519	151.675					
	MAMMA1002843	76.343	31.051	107.479	18.190	24. 282	30.456	19, 401	13.727
	MANMA 1002844	311.853	139. 150	228.560	66.881	72.282	201.758	152.946	94. 166
	MAMMA1002845	4.464	5.631	16.258	13.028	3.642	8.306	5.338	22.843
				10.230			178. 228	129, 737	278.807
	MAMMA1002857	77.604	209.913		167.148	50.200			
	MANMA 1002858	113.809	319.730	662.654	523.500	84. 144	532.413	382.518	1000.090
	MANSA 1002863	108. 297	33, 190	66, 980	38.305	26.112	45.735	86.883	51.987
					92.062	91.774	46.567	38, 439	58, 468
	MANMA 1002858	65. 375	102.643	253.035			40.301		10.400
	HAMMA1002869	85.453	22. 923	80.058	19.164	22.933	26.217	42.600	
	MAEMA 1002871	28.097	6.998	5.660	1.623	3.087	7,477	5.467	3.406
	HAMA1002875		16.542	18, 160	22. 528	23.110	21.099	24, 952	32, 949
		20. 954							
	MANNA 1002879	33. 352	14.773	9.446	6.359	8.506	13.275	30.077	23. 108
	MANWA 1002880	46.288	35.830	71.009	12.119	12.813	15.447	20, 107	22. 354
	MANA 1002881	57. 225	55, 154	238.977	25.333	27.378	18.964	34.053	52, 410
	MANMA 1002885	87. 039	28. 425	35. 323	14.016	29.952	34, 101	61.975	26. 271
	MANMA 1002886	398. 174	39.003	88.206	52.831	26.325	197.562	39.216	20. 561
	MACCA 1002887	45. 505	7.809	7.548	7.024	9.968	8, 271	13.675	5, 111
	MAYMA 1002890	65. 426	61.707	153.034	36.444	19.739	40.974	38, 649	41.029
	MAMMA 1002892	58, 445	53. 672	210.646	36.086	31.508	36, 186	13.729	35.746
	MAPMA1002893	76. 469	18.593	25.600	5.864	9, 192	24.826	20. 585	11.290
									21.933
	MAPMA 1002895	33.029	30.313	81.623	21.896	10.209	8. 431	11.614	1 (1.333

# Table 61

1APMA1002898	88. 538	24. 524	42. 725	9, 653	16.551	32, 137	42, 359	30, 615
AMMA1002905	191, 445	39.095	72.714	28, 234	32.209	91.200	60. 899	51.358
1AL 1002906	92.692	27.862	53. 273	26, 259	34, 130	57, 141	67, 635	26. 917
1AL 1002308	77.656	66. 964	209. 054	54.014	54, 429	43.639	58. 626	50. 901
					89.304	83.884	61.550	89.879
1AP1A1002909	157. 128	123.626	654. 652	152.777				
YAKYA1002918	55. 362	26. 201	35. 298	14. 931	10.960	19. 166	27.775	29. 119
MACMA1002925	50. 571	70.116	54. 395	18.071	27.814	43. 511	11.984	57.467
MACMA1002926	105. 041	221.644	119.112	66.217	73.866		1218.974	550. 265
MAKMA1002930	68.089	38.713	147.112	32.243	19, 181	31.875	24. 698	46.379
MAKMA1002937	207.866	61.711	89.764	38. 377	38, 050	97.677	156.876	119, 279
MAPMA1002938	34. 139	13.727	21.350	7.309	10.152	15. 165	14. 230	14. 534
HAMMA1002941	18.884	30. 845	50.805	19.591	7.699	16. 322	11. 528	24. 529
HAEMA1002947	63.095	31. 441	46.623	20.590	18.624	28. 594	29. 987	39.586
MAMMA1002964	43. 981	37.785	133. 836	22. 173	11.661	25. 346	15. 389	28. 296
MAMMA1002967	37.974	16,689	23. 126	13. 527	10.863	35.085	22.091	25.886
MALMA1002970	178. 268	124. 368	533. 590	120. 984	97.317	92.795	66.069	109,854
MAMMA1002971	99. 466	79.461	50.710	19.662	15, 091	40.745	37. 592	51.546
MAMMA1002972	83. 922	33. 377	50. 911	16.436	12.354	42.113	50. 137	45.819
MAPMA1002973	117.540	70.913,	318.513	45. 601	38.568	34.070	22.903	68.699
MAMMA1002979	80.771	204. 398	227.280	56.459	375, 745	119.386	122.750	226.538
MANYA 1002982	19. 895	9. 493	14. 202	6.265	0,000	0,000	0,000	5.076
MAPMA1002987	65. 397	50.918	156.507	28.534	30, 958	22.630	16. 594	36.952
MANMA1003003	104. 891	69, 630	125. 933	48.800	36.915	48.025	45.716	47.346
MALEMA 1003004	41. 353	106.059		111.746	92.691	59. 597	33.719	77.654
MAN-MA1003007	20. 423	21, 289	75.498	16.044	8, 909	15.878	6, 947	15, 193
NAMMA 1003011	45.615	37.641	29.754	23.843	21.157	33. 395	48. 907	39. 054
MA/MA1003013	65. 088	58. 284	49. 438	27.289	18, 877	31.768	67.950	59.419
MALMA1003015	36.817	29. 585	89. 251	19.826	4. 679	16, 602	6, 959	10, 432
MAMMA1003019	10.026	30. 107	5. 244	7, 467	2.375	6.403	3. 225	6. 184
MAMA1003020	48.046	31.761	50.515	13.842	17, 142	19.341	28. 497	20. 218
MAMA1003026	28.646	14. 274	3.514	8.603	6.618	9, 838	11.161	6. 781
MANA1003031	248. 219	140. 526	311.997	98, 494	105, 194	112.752	66, 462	132.570
MANNA 1003033	47.072	27. 208	130.132	44.811	42.096	33.806	17, 555	36.757
MANMA 1003035	102. 528	49.560	45. 025	30. 912	25. 924	64.046	42.175	56.246
MAMA1003039	37. 382	19.822	98.219	37. 555	17.115	27. 935	9. 656	25. 906
MARMA 1003040	76.014	95. 416		114, 795	84. 250	59. 989	42, 107	100.448
MARMA1003040	79.444	46.915	90.545	40.709	21. 121	25. 258	13.745	23.44
MAMMA 1003047	376. 340	121, 483	150, 100	91.015	100. 397	158. 521	175. 219	122. 400
MANUA 1003049	26, 899	9.631	9. 169	2, 907	5, 679	12.149	5.016	10.00
	38, 639	24.977		21, 811	15. 758	11.937	6.277	20.03
MANAGA 1003055	31, 238	13.811	76.695 32.121	15. 345	7.891	17.689	3, 176	18.14
MARMA 1003056		35. 596			19.335	28. 373	32.521	36.63
MAMMA 1003057	68. 258 43. 837	46, 015	34.053 117.875	23.862 31.178	11.361	17,068	9. 179	35.83
					3.215	6. 507	2.433	11.804
MAMMA1003075	16. 366	6.334 51.500	32.629 220.715	10. 374 36. 189	24.057	14. 625	14. 530	41.85
MAMMA 1003089	49, 867 22, 129	73, 102	15.615	27.304	11.693	9, 575	15, 986	84.96
	8. 240	37.313	24.078	8.354	10, 123	9, 662	24.609	12.39
MAMMA 1003095	44, 094	27.545		16.050	12. 184	15. 519	4. 930	23.72
MANYA 1003099 MANYA 1003102		18, 730	96.117	14.500	22.389	16. 929	20, 089	20.89
CMERCH LUDYINA	44, 491		31.447		10.720	11, 459	11.385	18.99
	25 077			14, 588	i iu. / ZV	1 11.435		
MANMA 1003 104	35. 977	19.146					17 722	72 21
MARMA1003104 MARMA1003113	41.697	21.092	30, 337	15.635	14.764	14, 690	17.723	<del></del>
LIANSA 1003104 LIANSA 1003113 LIANSA 1003126	41.697 20.042	21.092 39.595	30, 337 102, 916	15.635 21.241	14.764 15.167	14, 690 17, 921	20.876	26.56
HAMMA1003104 HAMMA1003113 HAMMA1003126 HAMMA1003127	41.697 20.042 57.961	21.092 39.595 27.221	30, 337 102, 916 102, 332	15. 635 21. 241 12. 486	14.764 15.167 8.002	14, 690 17, 921 12, 295	20.876 13.773	26.56 22.28
HAMMA1003104 HAMMA1003113 HAMMA1003126 HAMMA1003127 HAMMA1003131	41.697 20.042 57.961 267.516	21.092 39.595 27.221 37.924	30, 337 102, 916 102, 332 129, 263	15.635 21.241 12.486 66.563	14.764 15.167 8.002 86.667	14, 690 17, 921 12, 295 135, 209	20.876 13.773 95.293	26.56 22.28 83.25
HAMMATOO3104 HAMMATOO3113 HAMMATOO3126 HAMMATOO3127 HAMMATOO3131 MAMMATOO3131	41.697 20.042 57.961 267.516 22.855	21.092 39.595 27.221 37.924 14.308	30, 337 102, 916 102, 332 129, 263 5, 624	15.635 21.241 12.486 66.563 7.938	14.764 15.167 8.002 86.667 2.690	14. 690 17. 921 12. 295 135. 209 14. 984	20.876 13.773 95.293 7.633	26.56 22.28 83.25 17.26
HAMATOO3104 HAMATOO3113 HAMATOO3126 HAMATOO3127 HAMATOO3131 MAMATOO3131 MAMATOO3136	41. 697 20. 042 57. 961 267. 516 22. 855 6. 575	21. 092 39. 595 27. 221 37. 924 14. 308 9. 140	30, 337 102, 916 102, 332 129, 263 5, 624 33, 040	15. 635 21. 241 12. 486 66. 563 7. 938 4. 487	14.764 15.167 8.002 86.667 2.690 0.895	14. 690 17. 921 12. 295 135. 209 14. 984 1. 900	20.876 13.773 95.293 7.633 5.064	26. 56 22. 28 83. 25 17. 26 5. 31
LAMATOO3104 HAMATOO3113 HAMATOO3126 HAMATOO3127 HAMATOO3131 PAMATOO3131 HAMATOO3136 HAMATOO3136 HAMATOO3136	41.697 20.042 57.961 267.516 22.855 6.575	21.092 39.595 27.221 37.924 14.308 9.140 18.018	30, 337 102, 916 102, 332 129, 263 5, 624 33, 040 18, 562	15. 635 21. 241 12. 486 66. 563 7. 938 4. 487 11. 213	14.764 15.167 8.002 86.667 2.690 0.895 11.461	14.690 17.921 12.295 135.209 14.984 1.900 16.500	20.876 13.773 95.293 7.633 5.064 8.591	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81
LAMMATOO3104 HAMMATOO3113 HAMMATOO3126 HAMMATOO3127 HAMMATOO3131 HAMMATOO3131 HAMMATOO3136 HAMMATOO3146 HAMMATOO3150	41.697 20.042 57.961 267.516 22.855 6.575 14.105 311.806	21.092 39.595 27.221 37.924 14.308 9.140 18.018 87.992	30, 337 102, 916 102, 332 129, 263 5, 624 33, 040 18, 562 58, 938	15. 635 21. 241 12. 486 66. 563 7. 938 4. 487 11. 213 77. 271	14.764 15.167 8.002 86.667 2.690 0.895 11.461 104.739	14.690 17.921 12.295 135.209 14.984 1.900 16.500	20.876 13.773 95.293 7.633 5.064 8.591	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94
HAMMATOO3104 HAMMATOO3113 HAMMATOO3126 HAMMATOO3127 HAMMATOO3131 HAMMATOO3135 HAMMATOO3136 HAMMATOO3136 HAMMATOO3136 HAMMATOO3150 HAMMATOO3154	41, 697 20, 042 57, 961 267, 516 22, 855 6, 575 14, 105 311, 806 93, 002	21.092 39.595 27.221 37.924 14.308 9.140 18.018 87.992 39.912	30, 337 102, 916 102, 332 129, 263 5, 624 33, 040 18, 562 58, 938 37, 471	15. 635 21. 241 12. 486 66. 563 7. 938 4. 487 11. 213 77. 271 22. 819	14.764 15.167 8.002 86.667 2.690 0.895 11.461 104.739	14.690 17.921 12.295 135.209 14.984 1.900 16.500 165.139 31.742	20.876 13.773 95.293 7.633 5.064 8.591 115.042 26.299	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94 27. 56
HAMMA1003104 HAMMA1003113 HAMMA1003126 HAMMA1003127 HAMMA1003131 HAMMA1003131 HAMMA1003140 HAMMA1003146 HAMMA1003155 HAMMA1003155	41. 697 20. 042 57. 961 267. 516 22. 855 6. 575 14. 105 311. 806 93. 002 41. 709	21. 092 39. 595 27. 221 37. 924 14. 308 9. 140 18. 018 87. 992 39. 912 26. 308	30. 337 102. 916 102. 332 129. 263 5. 624 33. 040 18. 562 58. 938 37. 471 36. 508	15.635 21.241 12.486 66.563 7.938 4.487 11.213 77.271 22.819 14.326	14. 764 15. 167 8. 002 86. 667 2. 690 0. 895 11. 461 104. 739 19. 655 18. 674	14, 690 17, 921 12, 295 135, 209 14, 984 1, 900 16, 500 165, 139 31, 742 30, 842	20.876 13.773 95.293 7.633 5.064 8.591 115.042 26.299 23.489	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94 27. 56 18. 04
HAMMATOO3104 HAMMATOO3113 HAMMATOO3126 HAMMATOO3127 HAMMATOO3131 HAMMATOO3135 HAMMATOO3136 HAMMATOO3136 HAMMATOO3136 HAMMATOO3150 HAMMATOO3154	41, 697 20, 042 57, 961 267, 516 22, 855 6, 575 14, 105 311, 806 93, 002	21. 092 39. 595 27. 221 37. 924 14. 308 9. 140 18. 018 87. 992 39. 912 26. 308 32. 317	30, 337 102, 916 102, 332 129, 263 5, 624 33, 040 18, 562 58, 938 37, 471	15.635 21.241 12.486 66.563 7.938 4.487 11.213 77.271 22.819 14.326 12.108	14. 764 15. 167 8. 002 86. 667 2. 690 0. 895 11. 461 104. 739 19. 655 18. 674 24. 093	14.690 17.921 12.295 135.209 14.984 1.900 165.500 165.139 31.742 30.842 12.999	20.876 13.773 95.293 7.633 5.064 8.591 115.042 26.299 23.489 8.766	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94 27. 56 18. 04
HAMMA1003104 HAMMA1003113 HAMMA1003126 HAMMA1003127 HAMMA1003131 HAMMA1003131 HAMMA1003140 HAMMA1003146 HAMMA1003155 HAMMA1003155	41. 697 20. 042 57. 961 267. 516 22. 855 6. 575 14. 105 311. 806 93. 002 41. 709	21. 092 39. 595 27. 221 37. 924 14. 308 9. 140 18. 018 87. 992 39. 912 26. 308 32. 317 25. 338	30. 337 102. 916 102. 332 129. 263 5. 624 33. 040 18. 562 58. 938 37. 471 36. 508 147. 845 29. 052	15.635 21.241 12.486 66.563 7.938 4.487 11.213 77.271 22.819 14.326 12.108 18.551	14. 764 15. 167 8. 002 86. 667 2. 690 0. 895 11. 461 104. 739 19. 655 18. 674 24. 093 20. 826	14.690 17.921 12.295 135.209 14.984 1.900 165.500 165.139 31.742 30.842 12.999 32.639	20. 876 13. 773 95. 293 7. 633 5. 064 8. 591 115. 042 26. 299 23. 489 8. 766 35. 893	26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94 27. 56 18. 04 19. 93
HAMMA1003104 HAMMA1003113 HAMMA1003126 HAMMA1003127 HAMMA1003131 HAMMA1003131 HAMMA1003140 HAMMA1003146 HAMMA1003150 HAMMA1003155 HAMMA1003155	41. 697 20. 042 57. 961 267. 516 22. 855 6. 575 14. 105 311. 806 93. 002 41. 709 34. 876	21. 092 39. 595 27. 221 37. 924 14. 308 9. 140 18. 018 87. 992 39. 912 26. 308 32. 317	30. 337 102. 916 102. 332 129. 263 5. 624 33. 040 18. 562 58. 938 37. 471 36. 508 147. 845	15.635 21.241 12.486 66.563 7.938 4.487 11.213 77.271 22.819 14.326 12.108	14. 764 15. 167 8. 002 86. 667 2. 690 0. 895 11. 461 104. 739 19. 655 18. 674 24. 093	14.690 17.921 12.295 135.209 14.984 1.900 165.500 165.139 31.742 30.842 12.999	20.876 13.773 95.293 7.633 5.064 8.591 115.042 26.299 23.489 8.766	23. 81 26. 56 22. 28 83. 25 17. 26 5. 31 9. 81 46. 94 27. 56 18. 04 19. 93 33. 74 20. 13 7. 64

Table 62

	NB9N31000010	31.105	17.113	26.284	14.271	7. 540	17.180	16.220	11.568
	NB9N31000016	63.431	16, 195	24.879	17.001	16.740	25. 216	14.845	17.364
5	иВ9И31000043	87. 438	35, 161	58.144	20.813	36.473	36.956	51.575	34.673
3	NB9N31000045	83. 399	109. 448	62.101	95.653	93.734	94.218	166.654	74. 328
	NB9N31000054	41.821	12.636	37.831	15.025	15. 265	18, 963	10.894	13. 189
	NB9N31000076	22.822	22.709	57. 320	14. 223	12.517	9.029	11.713	24.494
	NB9N31000086	31, 281	74. 504	22.661	29. 164	11.744	29, 951	13.909	30,012
	NT2RU1000001	11.595	9, 900	11.540	4.467	4.016	8.823	6.775	5.184
	NT2RH1000018	333. 185	68. 022	171.103	77.680	48.418	138, 131	122.906	79.595
10	NT2RU1000032	37. 506	9, 768	23.088	9. 453	13. 222	16, 128	22.911	12.495
	NT2RH1000035	185. 573	46.513	81.354	56.890	39. 846	82.885	74.450	52. 553
		185. 843		116.479	50.830	36.658	98. 591	49.882	54. 356
	NT2RH1000037		60. 878		104, 606	82.108	214. 282	139.766	101.078
	NT2RH1000039	228. 804	172.849	444.715		52.898	89. 445	80. 537	184.618
	NT2RM1000042	55. 479	102.774	112.292	145. 900		0,000	5. 227	0.000
15	NT2RH1000055	1.083	0.593	0.000	0.000	0. 252			102.504
75	NT2RM1000059	212.057	100. 267	173. 989	78. 130	50.792	143. 445	83.189	10. 555
	NT2RH1000062	11.755	9. 438	11. 334	1. 925	2.705	2. 434	25.015	
	NT2RM1000065	153. 505	42. 956	56, 248	29.740	66.820	67.974	42.112	65. 531
	NT2RH1000066	26. 794	6. 5394	7. 914	2.716	6.609	8. 275	11.533	13.605
	NT2RM1000071	42. 919	126.091	61.623	97.378	24.665	45.008	74.491	266. 252
	NT2RH1000080	12.803	1.714	1.023	4.022	2. 135	8.919	13.254	4. 329
20	NT2RM1000086	393.857	146. 368	283.360	100.835	117.874	205. 973	155.085	102.325
	NT2RH1000092	12. 949	18.015	4. 187	6.602	2.600	0.000	5. 579	17.636
	NT2RM1000118	0.000	0. 276	0.000	0.180	0.000	0.000	0.000	0.655
	NT2RH1000119	18.719	5. 828	9.051	5. 794	3.873	6.048	19.700	10.812
	NT2RM1000121	2. 231	0.000	7.566	3. 177	3.735	3. 309	1.697	3.614
	NY2841000122	309.647	84. 904	138. 129	58.379	75.966	213.166	141,553	57.569
25	NT2RM1000127	14. 133	3. 707	2. 380	2.322	3.743	4. 212	8.594	5.786
	NY2RM1000131	1.661	1. 269	0. 348	0.000	0.768	0.000	2.271	2.221
	NT2RM1000132	10.432	7.649	9, 599	3.479	7. 287	11.592	13.046	10.752
	NT2RH1000153	39.773	9. 302	10.314	3.465	4.419	11.775	17. 131	12.503
	NT2RM1000184	85.966	171.937	58.982	34. 486	22.674	51.668	129.969	177.417
	NT2RM1000188	2. 149	4.607	0.000	0.000	1.586	1. 226	3.974	7. 121
30	HT2RM1000187	29. 354	12.303	16.019	17.222	15.020	17. 175	15. 232	18.703
30	NT2#M1000199	16. 274	0.000	17.316	6.834	4.725	5. 212	8.917	6.720
	NT2RH1000213	17. 361	14.639	43.481	9. 904	8.998	12. 127	6.422	10.141
	NT2RM1000215	8.787	10.858	90.070	4.505	89.435	12. 158	6.380	7.453
	NT2RM1000218	0.000	10.196	7.239	2.227	1.452	4. 273	8.324	4. 445
	NY2811000224	35.730	65.418	0.000	47.537	20.172	44. 102	26.563	63.368
	NT2RM1000236	52.706	47.803	20.481	19.138	42.513	21.813	58.118	100.492
35	NT2RM1000242	0,000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
	NT2RM1000244	13.988	12.654	6.957	9.937	6.047	8. 026	8. 938	3.968
	NT2RM1000252	283.006	144. 306	358.324	169.383	149. 200	192.609	174. 288	239.093
	NY2RM1000256	284. 496	113.021	203.771	67.954	94. 270	152. 181	132. 435	150.452
	NY2RM1000257	8. 203	8.081	9.713	9.716	0.000	5. 002	7.893	7.694
	NT2811000260	548. 461	312.072	494.663	164.454	249. 491	313.672	232, 568	270.549
40	NT2RH1000259	9. 472	7.461	6.606	10.004	8.876	5. 844	16.818	6.933
	NT2RH1000271	8. 917	1.259	3.857	2.440	2.317	4. 289	4. 982	5.727
	NT2RM1000272	83. 425	97.598	29.246	80.462	22.650	25. 350	34. 266	157.515
	NT2RM1000273	27.031	19.960	21.872	11.127	5.201	25.896	29.976	17.270
	HT2811000274	42. 234	91.340	28.306	26.224	11.534	34.723	32.623	85. 440
					7.205	7.361	10.397	4. 200	10.119
	INTZRM1000280	14, 289	1 12. 359	1 21.912	1.203	1	10.337		
15	NT2R11000280	14. 289 8. 249	12.359	17.445	4.671	9. 099	9.454	2. 185	1.092
45	NT2RH1000295	8. 249	4.916	17. 445				2.185 6.206	14.221
45	NT 2RE1000295 NT 2RE1000300	8. 249 41. 252	4. 916 31. 172	17.445 62.474	4.671	9.099	9.454	2. 185	14. 221 321. 054
45	NT2RH1000295 NT2RH1000300 NT2RH1000304	8. 249 41. 252 130. 855	4. 916 31. 172 217. 805	17. 445 62. 474 133. 583	4.671 15.266 142.504	9. 099 6. 023	9. 454 14. 825	2.185 6.206	14.221
45	NT2RE1000295 NT2RE1000300 NT2RE1000304 NT2RE1000314	8. 249 41. 252 130. 855 255. 347	4. 916 31. 172 217. 805 113. 392	17.445 62.474 133.583 165.204	4.671 15.266	9. 099 6. 023 77. 271	9. 454 14. 825 155. 874	2. 185 6. 206 78. 198	14. 221 321. 054
45	NT2R11000295 NT2R11000300 NT2R11000304 NT2R11000314 NT2R11000318	8. 249 41. 252 130. 855 255. 347 4. 002	4. 916 31. 172 217. 805 113. 392 22. 985	17. 445 62. 474 133. 583 165. 204 8. 505	4.671 15.266 142.504 56.831 14.343	9. 099 6. 023 77. 271 114. 936	9. 454 14. 825 155. 874 189. 937	2. 185 6. 206 78. 198 108. 461	14. 221 321. 054 113. 313 25. 194 14. 353
	NT2RE1000295 NT2RE1000300 NT2RE1000304 NT2RE1000314 NT2RE1000318 NT2RE1000335	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048	17. 445 62. 474 133. 583 165. 204 8. 505 6. 881	4. 671 15. 266 142. 504 56. 831 14. 343 7. 482	9. 099 6. 023 77. 271 114. 936 0. 836	9. 454 14. 825 155. 874 189. 937 6. 124	2. 185 6. 206 78. 198 108. 461 14. 391	14. 221 321. 054 113. 313 25. 194
<b>45</b>	NT2RE1000295 NT2RE1000300 NT2RE1000304 NT2RE1000314 NT2RE1000318 NT2RE1000335 NT2RE1000341	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157 41. 219	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048 3. 681	17.445 62.474 133.583 165.204 8.505 6.881 1.562	4. 671 15. 266 142. 504 56. 831 14. 343 7. 482 0. 000	9. 099 6. 023 77. 271 114. 936 0. 836 5. 897	9. 454 14. 825 155. 874 189. 937 6. 124 3. 558	2. 185 6. 206 78. 198 108. 461 14. 391 14. 151	14.221 321.054 113.313 25.194 14.353 6.704 85.201
	NT2RE1000295 RTZRE1000300 RT2RE1000304 NT2RE1000314 NT2RE1000315 NTZRE1000335 NTZRE1000341 NTZRE1000350	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157 41. 219 302. 316	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048 3. 681 74. 071	17.445 62.474 133.583 165.204 8.505 6.881 1.562 106.873	4.671 15.266 142.504 56.831 14.343 7.482 0.000 34.040	9.099 6.023 77.271 114.936 0.836 5.897 0.000	9. 454 14. 825 155. 874 189. 937 6. 124 3. 558 10. 884	2.185 6.206 78.198 108.461 14.391 14.151 5.578	14.221 321.054 113.313 25.194 14.353 6.704 85.201
	NT2RE1000295 RTZRE1000300 RT2RE1000304 NT2RE1000314 NT2RE1000315 HTZRE1000335 HTZRE1000350 NT2RE1000350	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157 41. 219 302. 316 6. 027	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048 3. 681 74. 071 0. 000	17. 445 62. 474 133. 583 165. 204 8. 505 6. 881 1. 562 106. 873 0. 000	4, 671 15, 266 142, 504 56, 831 14, 343 7, 482 0, 000 34, 040 1, 807	9.099 6.023 77.271 114.936 0.836 5.897 0.000 61.895	9. 454 14. 825 155. 874 189. 937 6. 124 3. 558 10. 884 149. 078	2.185 6.206 78.198 108.461 14.391 14.151 5.578 112.517	14.221 321.054 113.313 25.194 14.353 6.704 85.201
	NT2RE1000295 NT2RE1000300 NT2RE1000304 NT2RE1000314 NT2RE1000318 NT2RE1000331 NT2RE100034 NT2RE1000350 NT2RE1000354 NT2RE1000355	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157 41. 219 302. 316 6. 027 74. 362	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048 3. 681 74. 071 0. 000 158. 811	17. 445 62. 474 133. 583 165. 204 8. 505 6. 881 1. 562 106. 873 0. 000 209. \$78	4, 671 15, 266 142, 504 56, 831 14, 343 7, 482 0, 000 34, 040 1, 807 39, 101	9. 099 6. 023 77. 271 114. 936 0. 836 5. 897 0. 000 61. 895 0. 000	9. 454 14. 825 155. 874 189. 937 6. 124 3. 558 10. 884 149. 078 0. 921 249. 368	2.185 6.206 78.198 108.461 14.391 14.151 5.578 112.517 2.303 14.695	14.221 321.054 113.313 25.194 14.353 6.704 85.201
	NT2RE1000295 RTZRE1000300 RT2RE1000304 NT2RE1000314 NT2RE1000315 HTZRE1000335 HTZRE1000350 NT2RE1000350	8. 249 41. 252 130. 855 255. 347 4. 002 10. 157 41. 219 302. 316 6. 027	4. 916 31. 172 217. 805 113. 392 22. 985 10. 048 3. 681 74. 071 0. 000	17. 445 62. 474 133. 583 165. 204 8. 505 6. 881 1. 562 106. 873 0. 000 209. \$78	4, 671 15, 266 142, 504 56, 831 14, 343 7, 482 0, 000 34, 040 1, 807	9.099 6.023 77.271 114.936 0.836 5.897 0.000 61.895 0.000	9. 454 14. 825 155. 874 189. 937 6. 124 3. 558 10. 884 149. 078 0. 921	2.185 6.206 78.198 108.461 14.391 14.151 5.578 112.517 2.303 14.695 4.356	14. 221 321. 054 113. 313 25. 194 14. 353 6. 704 85. 201 1. 256 225. 724

Table 63

	NTODULANATA	02 502	0 616	40.007	20 751 1	21 004	C1 670 T	30 1/7	31 734
	NT2RH1000372	93. 583	9.616	49.097	28.751	33.904	61.678	39, 147	31.524
	NT2RH1000377	42.186	17.871	22.783	12.668	13. 142	15. 587	18.377	23.602
5	NT2R31000388	8.811	19. 351	1.155	5. 242	0.780	5. 795	6. 201	11.464
	NT2RH1000394	0.899	1.862	0.813	1.925	0.438	0.000	0.000	0.000
	HT2RH1000399	1.641	5. 386	0.000	2. 270	0.570	0.319	2.023	1.257
	NT2RM1000407	69.180	19.536	39.379	6. 299	21. 106	27. 229	14. 102	13.378
	NT2RH1000421	0.890	0.000	0.000	0.000	0. 456	0.150	0.000	0.000
	NT2RM1000422	102.028	152, 115	200.732	297.482	65. 137	134, 344	50. 452	241.878
10	NT2RH1000430	16.769	3. 286	12.402	4. 398	4. 506	12.149	11.238	7.508
10	NT2RM1000462	167.815	117.695	165.008	62.828	65. 795	81.561	72.026	118.786
	NT2RH1000499	16.037	22.127	75. 152	12.507	7.415	7. 335	41.299	22.217
	NT2RM1000512	126.610	24. 122	12.786	25.082	11.161	46.878	21.802	31.090
	NT2RM1000519	7.852	28.718	9.178	14.716	6.756	27.934	11.081	10.474
	NY 2RM 1000527	29.692	15. 338	24. 471	17.418	45. 221	59. 291	31. 450	14.020
	NT2RH1000539	14.790	19. 300	31.135	14.824	2. 560	6.669	3.751	10.774
15	NT2RM1000542	118.560	38. 555	21.020	20.675	29.849	30.176	22. 378	32. 507
	NT2RM1000553	37. 329	18.841	47.329	24. 533	23.901	33.590	34.084	33.966
	NT2RM1000555	77.352	46. 168	43.953	21.772	15.838	16.936	12.057	35.840
	NT2RM1000558	55.132	15. 424	20.508	7.987	7.249	8.885	23.984	21.919
	NT2RM1000563	39.161	14.058	17.872	12.234	8.871	14. 324	12.341	13.452
	NT2RM1000566	3.172	7.323	0.000	2.755	1.243	3, 584	2.944	4.754
20	NT2RM1000570	65. 428	72.508	44. 124	24. 498	15. 164	26.341	21.720	56.340
	HT2RH1000571	20.300	15.881	9.841	14. 197	7.525	7. 964	16.668	9.893
	NY2RH1000574	45. 305	32.953	5.746	5.977	1. 945	5.060	1.526	3.809
	NY2RH1000580	10.540	9. 295	12, 139	8.734	2.114	6. 532	5.687	7.120
	NT2RY1000620	11.778	12.782	21.632	15.504	5. 894	4. 488	3.359	17.303
	NT2RH1000523	3.914	2.515	0.416	3. 125	0. 251	0.715	0.355	2.159
25	NT2RM1000630	17.633	6.091	6. 532	3.910	2.095	8. 257	7.963	6.411
20	NT2RH1000633	5.563	70.230	93.799	22.316	42.967	24.174	6.091	43.328
	NY2RH1000534	3.427	3.869	2.248	1.997	0.487	0.000	1.258	3.039
	MT2RU1000642	87.902	31.353	26.846	11.421	21.495	75.074	66. 152	42.393
	NT2RU1000647	45, 410	65.742	56.619	55. 351	49, 439	30. 233	26.128	50. 923
	NT2RH1000648	25. 285	9.969	8.914	5. 538	3.383	6.085	5.045	5. 201
22	HT2801000650	22.370	15.864	19.881	11.036	29.031	8. 360	13,836	11.166
30	NT2RM1000661	23, 325	6, 294	12.692	7. 551	6.360	11.076	18.036	9.158
	NT2RM1000566	13.966	1.244	3. 221	1.629	1.543	4. 997	1.079	2.418
	NT2801000569	7.339	9.184	2.145	1.453	1.159	1.973	0.824	6.789
	NT2RE1000672	58.162	25.532	15.778	9, 171	22.446	58. 987	16.791	14.945
	NT2RM1000681	21.724	106.663	3.979	14.842	2.185	20. 284	16.034	21.688
	NYZREICOUS 91	4. 381	9. 202	2.832	3.483	1.268	0.878	2. 181	3.652
35	NT2RH1000698	31.943	17.379	9.609	16.495	5. 185	8.614	8.628	12.092
	NT2RH1000699	10.439	2.722	5. 406	4.115	3. 535	6. 367	10.784	8.214
	NT2RD1000702	32.110	7.097	17. 438	3.946	5.019	19.783	16.192	9.778
	NT2RM1000703	32.168	17.962	20.468	14.964	19, 912	19, 805	20. 940	16.286
	NT2RH1066704	25. 926	35.690	22. 230	11.998	15.536	38.075	52.384	26.689
	NT2R11000725	12.567	91,681	3.742	10.735	0. 262	10.694	14.773	17.602
40	NT2RH1000726	7.525	9.354	5. 608	7.297	2. 528	3. 884	3.237	8. 489
	NT2RM1000731	144.609	19.850	46.338	14.141	85.767	40. 231	32.791	30.972
	NT2RH1000741	14. 291	4.715	6. 122	2.576	3. 554	8. 230	5. 265	7.328
	NT2RU1000742	30.801	9. 241	6. 240	6.116	3. 655	11.131	7.680	11.315
	NT2RU1000744	69.419	21.887	27. 283	15.799	11.433	38. 093	24. 162	24. 347
	NT2RH1000746	12.863	7.631	12.042	6.326	6.665	9. 321	8.974	11.118
45	NT2RM1000747	24. 565	39.958	11.215	5. 537	1.866	7.009	10.940	21.461
	NY 2RD 1000752	13.148	7. 585	3. 359	5.748	4. 905	1.290	6.516	8.686
	HT2RU1000767	146.795	35. 621	33.719	11.495	31.430	63. 425	41.576	22.788
	MT2RD1000770	24. 395	7.712	21.569	11.954	11.449	9.412	14.053	17. 537
	HT2RU1000772	2.148	5. 100	1.271	2.181	0.000	1.505	6.132	3.034
	HT2RH1000779	284. 561	185.275	301.250	139.318	150.250	196. 541	146. 279	96.926
50	NT2811000780	9.227	9. 621	4. 260	6.864	3.591	4. 298	8.898	2.912
50	HT2RM1000781	0.000	0.000	4.468	0.666	2. 562	3.064	2.407	2.127
	NT2RD1000789	79.877	28.387	74. 545	23.140	28.956	35.852	51.230	46.548
	NY2RM1000800	4.947	10.706	34.906	3.517	6.856	4. 436	8.934	3.531
	NY2RD1000802	209, 372	41.025	60.767	12.693	69.721	155.310	133.291	27.049
	NY2R21000811	0.000	0.807	0.000	3.615	0.593	0.000	1.896	1.921

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. Table 64

NYTERIHODEZE   15,377   15,979   54,731   72,575   16,257   1,159   65,585   25,085   17,281000633   37,244   17,231   21,235   22,632   41,436   121,255   11,147   25,085   17,281000633   37,240   13,142   7,370   3,585   41,436   121,255   11,147   15,474   13,147   13,147   14		NT2RM1000826	55.971	29.000	28.733	20.800	13 355	7 100	20 144	22.706
							12.255	7. 195	28. 144	23. 708
	5	NIZALIOUUSI						121. 255		
	3			17.302	8.876	4.821	8.474	6.471	16.424	13, 119
					7.973	9.896	4.809	8.919	6.281	8, 562
			31.899	32. 922	28. 948	39.736	19.743	24, 819	26. 306	
		NT2RM1000848	10, 486		11.047					
		NT2RM1000850								
	10									
New Part										
No.			279. 225			198.222	264. 575		196. 436	
				223.032	628. 109	128.574	92.997	272, 161	183.324	165.845
No.		36, 148		71.518	26.137	22.828	37.610	46.674	48, 259	
Street		NT2RM1000874	94.766	25. 329	40.690	15.917	33,235	69.767	75, 898	
Bit   Part   1000845   12, 282   118, 317   233, 345   90, 226   109, 110   311, 111   130, 726   182, 827   187, 821   1000845   25, 895   146, 253   91, 557   73, 90, 807   63, 737   152, 93   156, 558   193, 445   187, 821   1000845   28, 474   12, 512, 13, 539   21, 037   13, 367   23, 959   22, 465   14, 058   187, 821   1000845   38, 928   11, 716   72, 313   31, 461   8, 537   188, 474   182, 264   80, 716   177, 821   1000845   39, 927   37, 943   30, 304   4, 018   6, 936   7, 286   6, 525   8, 715   177, 821   1000845   39, 927   37, 943   30, 304   4, 018   6, 936   7, 286   6, 525   8, 715   187, 821   1000810   13, 293   44, 90, 792   189, 446   46, 593   65, 444   122, 556   87, 145   72, 117   817, 821   1000810   13, 235   22, 507   15, 176   5, 355   3, 770   20, 204   15, 343   18, 656   187, 821   1000810   13, 244   90, 792   189, 446   46, 593   65, 444   122, 556   87, 145   72, 117   817, 821   1000924   29, 894   90, 792   189, 446   46, 593   65, 444   122, 556   87, 145   72, 117   817, 821   1000924   29, 895   12, 894   4, 90, 792   1, 784   10, 844   12, 256   87, 145   72, 117   817, 821   1000924   29, 895   12, 894   4, 946   27, 27, 79   20, 389   37, 281   20, 2	15	NY2RM1000882	32, 751							
Bir	NT2R91000883		118 317							
ST2R:1000893										
STERNIDOOSS   246, 338   100, 240   188, 653   51, 222   48, 537   188, 474   182, 264   80, 715										
B T2R 1000896   8, 028   11, 716   12, 431   3, 451   8, 055   10, 349   3, 262   6, 889										
BYT2RH1000895   20. 978   2. 796   3. 034   4. 018   6. 936   7. 286   6. 525   8. 715										
HITZREIIO00950	20									
STRENGOOST   12.235   22.507   15.176   6.355   3.770   20.204   15.343   18.656   17.2811000914   199.944   90.792   159.446   6.533   55.449   122.556   87.145   72.117   17.2811000913   36.141   16.161   19.116   13.229   8.831   18.002   10.279   10.389   17.2811000921   0.242   1.831   11.629   2.787   0.000   1.344   1.305   2.292   17.2811000922   29.895   12.894   4.946   4.788   7.984   10.844   15.108   57.49   17.2811000924   29.895   12.894   4.946   4.788   7.984   10.841   16.108   5.749   17.2811000927   34.940   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000925   13.349   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000925   13.349   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000925   13.349   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000925   13.349   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000925   13.349   11.379   12.531   13.272   6.919   7.215   10.132   8.882   17.2811000926   20.594   16.510   28.449   33.770   11.295   30.987   65.017   30.389   17.2811000927   69.241   15.184   3.909   6.863   4.421   9.222   33.371   10.246   8172811000927   69.241   51.581   16.390   19.560   15.357   27.890   33.675   45.410   17.2811000927   0.000	20									
STZRETIGOOSTI   199. 944   90. 792   169. 466   66. 693   65. 449   122. 556   87, 145   72. 117     STZRETIGOOSTI   36. 141   16. 161   19. 116   19. 127   10. 228   8. 831   18. 002   10. 279   10. 389     STZRETIGOOSTZ   13. 119   18. 060   5. 555   12. 140   3.037   3. 684   6. 526   16. 464     STZRETIGOOSTZ   92. 895   12. 894   4. 44   49. 44   48. 87   78. 48   6. 526   16. 464     STZRETIGOOSTZ   48. 046   34. 012   49. 155   23. 882   14. 687   14. 867   17. 603   20. 582     STZRETIGOOSTZ   48. 046   34. 012   49. 155   23. 882   14. 687   14. 867   17. 603   20. 582     STZRETIGOOSTZ   48. 046   34. 012   49. 155   23. 882   14. 687   14. 867   17. 603   20. 582     STZRETIGOOSST   33. 349   11. 379   12. 531   31. 272   6. 519   7. 215   10. 192   8. 882     STZRETIGOOSST   23. 347   14. 841   49. 930   16. 747   44. 584   52. 121   23. 270   34. 312     STZRETIGOOSSE   24. 574   14. 841   49. 930   16. 747   44. 584   52. 121   23. 270   34. 312     STZRETIGOOSST   20. 594   16. 610   28. 449   33. 770   11. 225   30. 897   65. 017   30. 389     STZRETIGOOSST   20. 594   16. 610   28. 449   33. 770   11. 225   30. 897   65. 017   30. 389     STZRETIGOOSST   20. 594   16. 561   28. 449   33. 770   11. 225   30. 897   65. 017   30. 389     STZRETIGOOSST   20. 594   51. 561   16. 390   5. 863   4. 421   8. 226   13. 337   10. 246     STZRETIGOOSST   20. 000   0.										46. 935
STZED1000913   36.141   16.161   19.116   13.229   8.891   13.002   10.279   10.389						6.355	3.770	20. 204	15, 343	
STZEN1000919   36.141   16.161   19.116   13.229   2.787   0.000   10.279   10.389   17.2811000922   31.119   18.060   5.555   12.140   3.037   3.684   6.526   15.464   17.2811000924   29.895   12.894   4.946   4.786   7.984   10.841   15.108   5.749   17.2811000927   48.046   34.072   49.155   23.882   14.687   17.603   20.582   17.2811000951   13.349   11.379   12.531   13.272   6.919   7.215   10.192   8.682   17.2811000956   5.337   16.522   6.739   2.246   6.192   6.379   6.215   8.675   17.2811000956   5.337   16.522   6.739   2.246   6.192   6.379   6.215   8.675   17.2811000956   5.347   14.841   49.930   16.747   44.584   52.121   23.270   34.312   17.2811000956   3.172811000956   3.185   49.930   16.747   44.584   52.121   23.270   34.312   17.2811000957   3.185   49.930   5.653   4.421   3.226   13.377   10.246   17.2811000957   3.594   16.510   28.449   33.770   11.295   30.987   65.017   30.389   17.2811000957   3.594   15.561   16.390   19.560   15.357   27.890   33.675   45.410   17.2811000978   0.000   0.			199. 944		169.446	46.693	65.449	122.556	87.145	72.117
ST2801000924			36. 141	16. 161	19.116	13. 229	8.891			
STZREI1000922   13.119   18.060   5.555   12.140   3.037   3.684   6.526   16.464     STZREI1000927   48.046   34.032   49.155   23.882   14.687   14.867   17.603   20.562     STZREI1000951   13.349   11.379   12.531   13.272   6.919   7.215   10.192   8.862     STZREI1000955   5.337   16.522   6.739   2.246   6.192   6.379   6.215   8.675     STZREI1000956   5.337   16.522   6.739   2.246   6.192   6.379   6.215   8.675     STZREI1000956   24.574   14.841   49.930   16.747   44.584   52.121   23.270   34.312     STZREI1000956   24.574   14.841   49.930   16.747   44.584   52.121   23.270   34.312     STZREI1000956   24.574   14.841   49.930   16.747   44.584   52.121   23.270   34.312     STZREI100957   69.241   51.561   16.390   19.560   15.357   27.890   33.675   45.410     STZREI1000957   69.241   51.561   16.390   19.560   15.357   27.890   33.675   45.410     STZREI1000958   7.275   2.308   2.120   2.059   1.138   1.293   1.746   4.769     STZREI1000952   7.275   2.308   2.120   2.059   1.138   1.293   1.746   4.769     STZREI1000952   7.275   2.308   2.120   2.059   1.138   1.293   1.746   4.769     STZREI1000958   13.759   6.798   22.345   7.467   6.192   11.494   1.942   9.039     STZREI1000958   12.087   15.19   14.959   10.866   9.132   2.303   4.549   41.654     STZREI1001002   46.253   5.707   19.271   15.499   18.065   33.283   21.225   33.831     STZREI1001003   4.107   33.647   23.710   23.835   3.391   10.638   6.307   14.681     STZREI1001103   2.532   6.694   3.303   6.673   8.650   15.822   23.168   23.128     STZREI1001107   8.644   4.934   1.241   2.455   1.873   2.894   4.062   7.068     STZREI1001108   224.654   234.771   124.092   88.774   75.070   85.777   124.713   84.612     STZREI1001108   224.654   234.771   124.092   88.774   75.070   85.777   124.713   84.612     STZREI1001076   31.69   2.991   1.316   0.000   0.352   2.727   2.878   3.632     STZREI1001076   31.69   2.991   1.316   0.000   0.352   2.777   2.778   3.643   3.647   3.344   3.920   3.647   3.643   3.647   3.647		NT2RM1000921	0. 242	1.831	11,629		0.000			
NTZRI1000924   29.895   12.894   4.946   4.788   7.984   10.841   16.108   5.749	25	NT2RM1000922								
STZENTOOO997										
STZRE1000951   13.349   11.379   12.531   13.272   6.919   7.215   10.192   8.882   STZRE1000956   5.337   16.522   6.739   2.246   6.192   6.379   6.215   8.675   STZRE1000950   24.574   41.841   49.930   16.1747   44.584   52.121   23.270   34.312   34.512   3										
STZENIBODSS										
NTZED1000960										
NTZREJ1000961   20.594   16.610   28.449   33.770   11.295   30.987   55.017   30.389   NTZREJ1000962   1.479   8.158   49.309   6.863   4.421   9.226   13.337   10.246   NTZREJ1000978   69.241   51.561   16.330   19.560   15.357   27.890   33.675   45.410   NTZREJ1000978   0.000   0										
NTZRM1000982										
NT2RH1000973	30									30. 389
ST2RH1000978   0.000   0.000   0.000   0.000   0.000   0.000   0.000   0.000   0.368   ST2RH1000982   7.275   2.308   2.120   2.059   1.138   1.293   1.746   4.769   ST2RH1000984   12.087   15.119   14.969   10.866   9.132   2.303   4.549   14.654   1.942   1.942   9.039   ST2RH1000084   12.087   15.119   14.969   10.866   9.132   2.303   4.549   14.654   1.942   9.039   ST2RH1001002   46.263   5.707   19.271   15.499   18.065   33.283   21.225   33.831   ST2RH1001008   4.937   4.656   0.740   4.466   2.544   3.192   3.215   10.971   ST2RH1001001   67.834   16.031   21.431   8.274   20.203   46.979   40.030   18.121   ST2RH100101   57.834   16.031   21.431   8.274   20.203   46.979   40.030   18.121   ST2RH100101   25.323   6.684   3.303   6.673   8.650   15.882   23.168   23.126   ST2RH100101   224.654   234.771   124.092   68.774   75.070   85.777   124.713   184.612   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001026   23.653   12.510   10.387   14.301   5.568   12.341   14.518   17.008   ST2RH1001026   33.653   2.991   1.316   0.000   0.352   2.727   2.878   3.632   ST2RH1001036   3.679   5.544   0.768   1.254   0.973   4.81   1.761   5.391   ST2RH1001036   3.679   5.544   0.768   1.254   0.973   4.81   1.761   5.391   ST2RH1001032   3.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851   ST2RH1001032   3.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851   ST2RH1001032   3.245   3.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704   ST2RH1001032   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799   ST2RH1001032   63.705   50.432   105.417   34.113   20.331   17.230   16.37								9. 226	13. 337	10. 246
NT2RH1000982			69. 241	51.561	16.390	19. 560	15. 357	27.890	33. 675	45. 410
ST2RE100982   7. 275   2. 308   2. 120   2. 059   1. 138   1. 293   1. 746   4. 769     ST2RE1000981   13. 759   6. 798   22. 345   7. 467   6. 192   11. 494   1. 942   9. 039     ST2RE1001002   45. 263   5. 707   19. 271   15. 499   18. 065   31. 22   2. 303   4. 549   14. 654     ST2RE1001003   14. 107   33. 647   23. 710   23. 835   3. 391   10. 638   8. 307   14. 681     ST2RE1001003   4. 937   4. 696   0. 740   4. 466   2. 544   3. 192   3. 215   10. 971     ST2RE1001011   67. 834   16. 031   21. 431   8. 274   20. 203   46. 979   40. 030   18. 121     ST2RE1001013   25. 323   6. 694   3. 303   6. 673   8. 650   15. 882   23. 168   23. 126     ST2RE1001010   224. 654   234. 771   124. 092   68. 774   75. 070   85. 777   124. 713   184. 612     ST2RE1001016   224. 654   234. 771   124. 092   68. 774   75. 070   85. 777   124. 713   184. 612     ST2RE1001026   23. 853   12. 510   10. 387   14. 301   5. 568   12. 341   14. 618   17. 008     ST2RE1001026   21. 614   13. 830   4. 261   8. 481   4. 770   7. 687   17. 274   10. 663     ST2RE1001036   3. 169   2. 991   1. 316   0. 000   0. 352   2. 727   2. 878   3. 632     ST2RE1001056   3. 169   2. 991   1. 316   0. 000   0. 352   2. 727   2. 878   3. 632     ST2RE1001056   3. 169   2. 991   1. 316   0. 000   0. 1348   1. 228   3. 011     ST2RE1001074   32. 455   14. 324   28. 723   10. 090   6. 573   10. 841   7. 837   10. 538     ST2RE1001076   7. 339   4. 891   0. 792   2. 511   0. 000   5. 644   6. 602   2. 026     ST2RE1001076   7. 339   4. 891   0. 792   2. 511   0. 000   5. 644   6. 602   2. 026     ST2RE1001076   3. 921   7. 236   4. 420   3. 000   0. 000   0. 564   6. 602   2. 026     ST2RE1001076   7. 339   4. 891   0. 792   2. 511   0. 000   5. 644   6. 602   2. 026     ST2RE1001076   7. 339   4. 891   0. 792   2. 511   0. 000   5. 644   6. 602   2. 026     ST2RE1001076   7. 339   4. 891   0. 792   2. 511   0. 000   5. 644   6. 602   2. 026     ST2RE1001082   61. 133   28. 559   80. 293   36. 442   13. 840   23. 671   15. 948   30. 844     ST2RE1001			0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.368
STRENGOO991   13.759   6.798   22.345   7.467   6.192   11.494   1.942   9.039     NT2RHIGOO994   12.087   15.119   14.969   10.866   9.132   2.303   4.549   14.654     NT2RHIGOTOO3   14.107   33.647   23.710   23.835   33.931   10.638   8.307   14.681     NT2RHIGOTOO3   14.107   33.647   23.710   23.835   33.931   10.638   8.307   14.681     NT2RHIGOTOO1   67.834   16.031   21.431   8.274   20.203   46.979   40.030   18.121     NT2RHIGOTOTI   25.323   6.694   3.303   6.673   8.650   15.882   23.168   23.126     NT2RHIGOTOTI   8.644   4.934   1.214   2.455   1.873   2.894   4.062   7.068     NT2RHIGOTOTI   8.644   4.934   1.214   2.455   1.873   2.894   4.062   7.068     NT2RHIGOTOTI   224.654   234.771   124.092   68.774   75.070   85.777   124.713   184.612     NT2RHIGOTOZE   23.853   12.510   10.387   14.301   5.568   12.341   14.618   17.008     NT2RHIGOTOZE   23.853   12.510   10.387   14.301   5.568   12.341   14.618   17.008     NT2RHIGOTOZE   11.717   13.271   17.437   18.862   5.641   12.231   8.930   11.443     NT2RHIGOTOZE   11.717   13.271   17.437   18.862   5.641   12.231   8.930   11.443     NT2RHIGOTOZE   3.164   13.830   4.261   8.481   4.770   7.687   17.274   10.663     NTZRHIGOTOZE   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NTZRHIGOTOZE   3.0879   5.544   0.768   1.254   0.973   4.181   1.761   5.391     NTZRHIGOTOZE   3.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NTZRHIGOTOZE   63.705   50.432   105.417   34.113   20.301   17.852   6345   2.505   12.387     NTZRHIGOTOZE   63.705   50.432   105.417   34.		NT2RM1000982	7. 275	2.308	2, 120	2.059	1, 138	1, 293	1.746	
NT   REM   1000   12   12   13   14   14   15   10   18   16   10   18   18   10   10   18   18   19   18   18   19   18   18		MT2RM1000991	13.759	6.798						
ST2RE1001002	35	MT2RM1000994	12.087							
NTZRM1001003	55	NT2RH1001002								
NT2RM1001008										
NT2RM1001011   67.834   16.031   21.431   8.274   20.203   46.979   40.030   18.121										
### RETERMIQUIDIS  ### 25.323										
### RT2RM1001017										
ST2RH1001018   224.654   234.771   124.092   68.774   75.070   85.777   124.713   184.612   ST2RH1001026   23.853   12.510   10.387   14.301   5.568   12.341   14.618   17.008   ST2RH1001028   11.717   13.271   17.437   18.862   5.641   12.231   8.930   11.443   ST2RH1001043   21.614   13.830   4.261   8.481   4.770   7.687   17.274   10.663   ST2RH1001044   21.983   20.272   44.315   8.181   4.171   5.809   4.623   9.566   ST2RH1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632   ST2RH1001063   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391   ST2RH1001056   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011   ST2RH1001072   13.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851   ST2RH1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538   ST2RH1001032   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799   ST2RH1001032   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799   ST2RH1001032   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844   ST2RH1001102   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2RH1001102   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.08										
ST2RM1001026   23.853   12.510   10.387   14.301   5.568   12.341   14.618   17.008	40									
NT2RM100102B   11.717   13.271   17.437   18.862   5.641   12.231   8.930   11.443     NT2RM1001043   21.614   13.830   4.261   8.481   4.770   7.687   17.274   10.663     NT2RM1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NT2RM1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NT2RM1001053   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391     NT2RM1001056   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011     NT2RM1001056   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011     NT2RM1001072   13.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851     NT2RM1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538     NT2RM1001066   7.339   4.891   0.792   2.511   0.000   5.644   6.602   2.026     NT2RM1001062   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RM1001065   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704     NT2RM1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RM1001102   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772     NT2RM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.156     NT2RM1001101   5.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089										
NT2RM1001043   21.614   13.830   4.261   8.481   4.770   7.687   17.274   10.663   NT2RM1001044   21.983   20.272   44.315   8.181   4.171   5.809   4.623   9.566   NT2RM1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632   NT2RM1001053   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391   NT2RM1001056   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011   NT2RM1001072   13.706   7.601   5.972   3.306   0.165   3.139   5.672   5.851   NT2RM1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538   NT2RM1001076   7.339   4.891   0.792   2.511   0.000   5.644   6.602   2.026   NT2RM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799   NT2RM1001085   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704   NT2RM1001082   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704   NT2RM1001082   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772   NT2RM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   NT2RM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   0.000   1.156   NT2RM1001102   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089										17.008
NT2RH1001044   21.983   20.272   44.315   8.181   4.171   5.809   4.623   9.566     NT2RH1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NT2RH1001063   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391     NT2RH1001065   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011     NT2RH1001072   13.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851     NT2RH1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538     NT2RH1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RH1001085   13.921   7.236   4.420   3.206   4.563   0.965   5.984   4.704     NT2RH1001082   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   7.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   7.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   7.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   7.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   3.980   7.852   6.345   2.505   12.387     NT2RH1001103   4.293   14.550   11.888   7.286   7.266   5.412   8.469   12.089								12.231		11.443
NT2RH1001059   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NT2RH1001063   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391     NT2RH1001056   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011     NY2RH1001072   13.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851     NT2RH1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538     NT2RH1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RH1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RH1001085   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704     NT2RH1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RH1001076   2.299   0.000   0.000   0.000   2.005   1.301   2.772     NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RH1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   0.000   1.155     NT2RH1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   0.000   1.155     NT2RH1001105   0.000   0.418   0.000   0.686   0.000   0.										
NT2RM1001089   3.169   2.991   1.316   0.000   0.352   2.727   2.878   3.632     NT2RM1001083   0.879   5.544   0.768   1.254   0.973   4.181   1.761   5.391     NT2RM1001086   3.011   3.061   0.000   3.241   0.000   1.348   1.228   3.011     NT2RM1001072   13.706   7.601   5.972   2.306   0.165   3.139   5.672   5.851     NT2RM1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538     NT2RM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RM1001085   13.921   7.236   4.420   3.206   4.563   0.965   5.984   4.704     NT2RM1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RM1001102   2.299   0.000   0.000   0.000   0.000   2.005   1.301   2.772     NT2RM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   0.000   1.155     NT2RM1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089			21.983	20. 272	44. 315	8, 181	4.171	5, 809	4, 623	9. 566
ST2RM1001063	45	NT2RM1001059	3. 169	2.991					2.878	3, 632
NY 2	40	NT 28M 100 1063	0.879	5, 544	0.768					
SYZEM1001072   13.706   7.801   5.972   2.306   0.165   3.139   5.672   5.851     SYZEM1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538     SYZEM1001066   7.339   4.891   0.792   2.511   0.000   5.644   6.602   2.026     SYZEM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     SYZEM1001095   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704     SYZEM1001095   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     SYZEM1001102   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772     SYZEM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     SYZEM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.155     SYZEM1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089		NT 2RM 1001086								
ST2EM1001074   32.455   14.324   28.723   10.090   6.573   10.841   7.837   10.538   ST2EM1001076   7.339   4.891   0.792   2.511   0.000   5.644   6.602   2.026   ST2EM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799   ST2EM1001085   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704   ST2EM1001082   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844   ST2EM1001082   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772   ST2EM100103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   ST2EM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.156   ST2EM1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089   1.2089										
ST2RM1001076										
NT2RM1001082   63.705   50.432   105.417   34.113   20.331   17.230   16.378   21.799     NT2RM1001085   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704     NT2RM1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RM1001102   2.299   0.000   0.000   0.000   0.000   2.005   1.301   2.772     NT2RM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.155     NT2RM1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089										
NT2RH1001095   13.921   7.236   4.420   3.206   4.563   0.966   5.984   4.704     NT2RH1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RH1001102   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772     NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RH1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.156     NT2RH1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089										
NT2RID1001092   16.133   28.559   80.293   36.442   13.840   23.671   15.948   30.844     NT2RID1001102   2.299   0.000   0.000   0.000   0.000   2.005   1.301   2.772     NT2RID1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387     NT2RID1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.155     NT2RID1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089	50									
NT2RM1001102   2.299   0.000   0.000   0.000   0.000   2.006   1.301   2.772   NT2RM1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   NT2RM1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.156   NT2RM1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089	•	MT250145								
NT2RH1001103   4.293   14.550   11.888   3.980   17.852   6.345   2.505   12.387   NT2RH1001105   0.000   0.418   0.000   0.686   0.000   0.000   0.000   1.156   NT2RH1001112   6.983   5.403   12.985   7.889   7.226   5.412   8.469   12.089										
NT2RH1001105 0.000 0.418 0.000 0.686 0.000 0.000 0.000 1.156 NT2RH1001112 6.983 5.403 12.985 7.889 7.226 5.412 8.469 12.089						0.000	0.000	2.006	1.301	2.772
NT2RU1001105 0.000 0.418 0.000 0.686 0.000 0.000 0.000 1.156 NT2RU1001112 6.983 5.403 12.985 7.889 7.226 5.412 8.469 12.089				14, 550	11.888	3. 980	17.852	6.345	2.505	12. 387
NTZRE1001112 6.983 5.403 12.985 7.889 7.226 5.412 8.469 12.089			0,000	0.418						
55		NT2RM1001112	6. 983							
	55									

Table 65

	NT2RM1001115	100.486	24.788	67.251	18.301	19. 421	53.304	29.318	21.097
	NT2RH1001122	18. 980	19.515	19.938	11,109	10.211	34. 308	33, 955	13.422
	NT2RU1001136	4.811	3.751	2.520	1, 126	0.765	2.194	2.817	5, 117
5	NT2RM1001139	78.791	18, 931	27,710	8. 382	21.060	31.349	14.028	14, 521
	NT2R112000003	27.773	13. 438-	12.296	3. 254	10. 288	4. 103	14.697	22.880
	NT2RH2000006	64, 154	36. 637	117, 073	30. 277	27. 783	25. 842	17.647	24. 349
	HT2RH2000010	57. 805	33.217	60.148	20.749	86.788	23.487	19.722	22.651
	NT2RM2000013	24.877	27. 244	40. 874	15, 590	40.045	30.831	48. 932	36.344
	HT2RM2000030	68.595	26. 308	27.271	17. 595	25. 608	41. 165	43. 837	27.939
10				59. 847	11.737	13.094	11.681	12. 137	11.426
	NT2R12000032	22. 984	13.418	28.101	23.568	9, 740	51.053	23.006	23. 405
	NT 2RM2000039	35.892	5.887 9.200	20.886	10.060	5. 098	11, 101	20, 459	10.744
	NT2RM2000042	7.936			5,779	5. 195	6. 720	11, 106	5.712
	NT2RM2000092	12.085	11.085 31.271	15.415 57.365	24.041	26. 832	24. 640	12.930	20.135
	NT2RM2000093	51.998				29. 563	48. 487	33. 388	54. 246
15	NT2RH2000101	34, 341	46. 587	54. 294	27. 692 33. 739	39.845	53. 753	69. 151	73.279
	NT2RH2000104	73.163	48. 315	58.786		11.012	23.770	21. 401	22, 254
	NT2RM2000124	35.818	16. 923	31.954	10.723 12.467	9. 797	13. 085	10.315	17, 050
	NT2RM2000155	31, 139	23.019	27. 033 87. 171	59. 579	62.006	74. 514	126. 950	91. 326
	NY2RM2000191	151.075	54.651		4. 582	1, 137	2. 242	1, 413	0.000
	NT2RM2000192	0.760	2, 690 '	0.971	31. 157	21. 570	60. 155	49.672	39. 127
20	NT 2 RM 2000 239	92.578	36.060 69.966	71.933 77.545	23. 453	53.412	78, 029	64. 223	83, 906
	NT2RH2000240	104, 218 70, 281	31.167	42.733	18.007	14. 544	13. 466	26. 176	42. 298
	NY2RD2000247	72.366	22. 586	52.512	23.631	19.076	29. 100	50.616	50. 848
	NT2RD2000250	90. 122	33.799	39, 931	17. 198	9. 865	44. 083	74.558	29. 086
	NT2RM2000250	340.036	40.469	141.962	35. 653	77.794	188. 072	216.739	59. 426
	HT2RM2000265	24, 506	4.177	38.440	1, 951	3. 495	14. 217	14. 995	14. 683
25	NY28112000287	131.692	88. 080	127. 535	51.611	38. 294	53. 574	55. 104	70.583
	NT2RM2000306	45. 342	24, 950	44.593	13.884	40. 471	40.133	22.666	33. 254
	NT2R12000312	13.383	57.043	78.915	13. 258	60.055	90. 975	183. 575	38. 391
	NT2RH2000322	33.318	18.077	22.354	11,030	6.002	8. 829	16. 952	15.344
	NT2RD2000343	70.618	78. 514	302.242	43, 179	64, 338	35.838	84, 150	77, 161
	NT2RM2000359	79. 203	25. 437	34. 945	19.556	16. 348	47. 922	31.041	20.663
30	NY2RD2000352	138, 367	75.052	100.195	73.363	49. 275	128.683	126.847	106. 528
	NT2RH2000363	41.249	17, 128	40.363	12.316	18.047	6. 982	11.907	9. 239
	HY28112000360	225. 366	121.451	100.718	49.727	89.663	128. 354	136.054	93. 203
	HT2RM2000371	88.897	208. 325	97.848	212.525	33.081	80. 287	140.890	131.756
	NY2RM2000374	54.398	55.656	153.004	34.316	25.750	36.072	34, 151	51.955
	HT2R112000387	31.537	35.012	44. 269	24. 245	23.611	19.094	24. 288	26.745
35	HY28H20003H3	43.873	18.662	32.917	12.496	14. 167	17.560	23. 452	33. 102
	NT2RM2000395	11.936	2.901	3. 145	1.722	4. 564	6. 102	4.725	9. 257
	NT2RM2000402	26.540	28.616	42.681	18.209	10.970	24.876	20.077	26. 993
	NT2HM2000405	29.390	26.302	56. 236	18.391	18. 524	17.673	19.408	19. 435
	RYZRUZ000407	213.973	77.583	145. 459	42.798	73.678	124. 360	103. 989	122.635
	NT2R02000410	46.375	23, 782	29. 096	10.711	13. 331	26.855	27. 992	20.820
40	NY28112000420	41,781	29. 100	39.676	24.872	16.605	26.730	29.136	43.708
	HY2RM2000422	400.274	145. 824	265. 042	51.828	73.571	186.812	131.563	125. 088
	HY2RM2000423	119,707	56. 563	272.757	58.213	50.981	60. 353	42. 529	86.903
	NT2R112000452	44.543	24.735	36.727	13.780	10.160	32. 134	23. 468	26.716
	HT2RH2000469	28.062	19.762	14. 685	5.603	7. 485	22.242	10.716	6. 249
	NT2R112000480	57.984	29.556	42.743	16.403	19.316	36. 503	21. 106	31.221
45	NT2R12000497	44.862	39, 966	107.651	23.488	15. 277	19. 316	13.374	15.412
	NY2RM2000502	49.184	33, 683	39.515	14. 256	18.792	23. 598	23.921	27. 778
	NY2R112000504	53.653	30.376	46. 453	19.836	22.267	39. 106	28. 508	19. 188
	MT2RM2000514	40.702	23. 938	23.980	9.704	12.601	20.319	19.147	14. 506
	NT2R12000522	6.782	0.000	4. 730	3.680	1.616	2.008 16.595	10.471	17.045
	NT2RM2000540	28. 543	24. 938	24. 326	8.984	9.799		0.000	0.000
50	NT2RD2000556	0.000	0.000	0.000	0.000	0.000	42.730	24. 240	28. 218
	NT28112000565	52. 454	32. 231	48. 697	17.373	14.758	32.508	18, 105	35. 032
	NY2R12000565	31.997	22. 486	34. 598	11.793	7.665	44. 727	22. 394	28.766
	IJT 2 R2 2000567	57.110	29. 153	45. 058	10.738	15.606	58. 576	40. 151	50.117
	NT2812000569	113.652	91.632	187. 867	40.645	36.420	83.608	36. 221	60. 695
	NT2R12000577	61.308	16.114	35. 195	12.694	14. 986	79. 582	62. 192	40.676
55	HT2R02000581	152.797	45. 271	66.363	20.096	32.397	1 13. 302	1 04. 136	1 40.010

Table 66

	NTARWARATOR	AC 163 I	03 700 1	104 000 1	45 000 1	45 777 1	67 766	EO 430 I	50. 631
	NT2RM2000582	96.163	83.789	104.868	37.893	45.777	67.766	50.428	
	NT2RM2000588	109.847	89. 480	119. 521	70.534	32. 168	143.491	88. 984	95. 908
_	NT2RM2000589	91.130	45. 398	66. 143	21.774	22.548	80.656	43.864	35. 379
5	NT2RM2000594	31.068	22.138	28.684	10.809	13. 325	34. 179	10.310	16.391
	NT2RH2000599	275. 423	132.063	221.911	86.738	66.363	237.294	209. 381	119.304
	NT2RM2000509	25.687	13.378	20.025	9.729	14. 321	19. 395	17.956	8. 545
	NT2RH2000612	40.704	19.012	36, 338	9.471	15. 531	27.049	24.872	30.269
	NT2RH2000622	45. 492	46. 307	46.012	27.097	17.426	48. 495	30.090	42.927
	NT2RH2000623	279.041	219.374	245, 200	90.410	123.723	286, 194		144. 950
10							29. 084	27.783	54. 409
. •	NT2RH2000624	52.551	88.174	87.665	60. 273	35. 044			
	NT2RM2000632	15.461	13.673	11.853	13.378	8. 044	7.114	6.910	5.808
	NT2RM2000635	24.726	21.442	42.243	17.900	14. 353	23. 119	10.306	20.675
	NT2RM2000636	45. 247	47.662	62.828	24.460	33. 311	28.868	35.751	35. 343
	NT2RM2000639	34. 707	19. 290	26. 594	15. 919	12.875	28. 297	20.526	11.317
	NT2RM2000649	39.662	37.102	62.088	31. 152	32. 252	42. 335	27.796	50.424
15	NT2RM2000658	53. 598	26.723	55. 360	19.176	26. 348	46.815	23.949	20.812
	NT2RM2000660	84. 441	62.193	66.364	13.329	36.417	48. 257	23.694	40.215
	NT2RM2000669	17. 352	23.877	38.180	11.181	16.885	17.594	13.008	20.479
	NT2RM2000689	118.126	102.565,	102.237	102.435	37.057	156. 147	96.539	140.413
	NT2RH2000691	29. 467	12.787	29.631	9. 783	15. 294	28. 392	15.401	17.161
	NT2RM2000714	238.396	61.067	122. 264	38. 290	60. 785	222. 914	188, 827	77.434
20						7, 922	8. 962	7.572	22.010
	NT2RM2000718	9.515	10.199	19.686	5. 036	29, 863	38. 537	30. 201	17.415
	NT2RM2000732	44. 022	24. 869	42.915	12. 209				59.237
	NT2RM2000735	112.208	47.966	111. 282	57. 228	38, 980	78. 590	45.888	
	MT2RM2000740	23. 990	62.438	143.286	24.030	26. 159	35. 449	22.001	29.845
	NT2RM2000743	15. 424	14. 901	23. 591	12.391	9. 779	16.339	8. 950	8.560
	NT2RM2000772	79.885	34. 020	54. 908	31.068	31.256	64.893	44. 735	55. 557
25	NT2RM2000773	56. 846	36.465	77. 155	26. 645	32. 523	60.130	42.946	53.958
	NY2RM2000776	56.550	40.820	69.793	43.736	22. 285	89. 348	33. 285	45.221
	NT2RM2000784	54. 586	33.888	45. 181	19.559	21. 292	43. 103	25. 540	42.124
	NT2RM2000795	169. 462	132.660	456.283	117.450	94. 702	91.566	59.832	91.914
	NT2RM2000796	12.942	12.033	20. 129	5.817	6.070	11.596	8.538	11.009
	NT20M2000798	67.292	147. 984	71.980	42.802	43. 127	85. 427	63.126	132.706
30	HT2RH2000801	145. 709	152.451	160.966	85. 365	73.827	214. 221	157. 384	174.371
	NT2R12000821	29.716	25. 994	36.976	14. 293	9. 638	63.476	12. 133	3.427
	NT2RM2000829	77. 695	36.834	148.015	32.077	69.569	70.012	26. 103	73.222
	NY2RM2000837	85.748	27.100	51.022	19.432	22, 405	48.733	36.614	45.277
						16, 935	130. 595	55. 870	42.226
	NT2R112000924	41.170	22.739	31.818	6. 582			28. 135	30.171
0.5	NT2RH2000930	45. 514	31.120	39.165	20.017	17. 433	49.111		
35	NT2RM2000937	85. 092	19.912	28.613	13.728	34. 425	55. 176	53.959	15.755
	NT2RM2000939	63. 956	41.986	59. 137	18.909	23.056	57.088	26.370	29.465
	NT2RM2000942	141. 275	345.015	119.378	242.434	78. 282	274.472	112.054	436.171
	NT2RH2000951	32. 383	20.717	12.763	17.041	10.179	32.704	19. 494	30.498
	NY2RM2000952	33. 160	18.882	34.052	15. 194	27.783	44. 540	16.881	31.012
	NT2RM2000966	54. 007	44. 546	57.551	30. 397	27.965	78.353	44. 947	77.916
40	NT2RM2000973	96.188	97.082	100.373	31.654	38.259	115.479	60.146	151.200
	NY2RM2000983	66.024	27.357	40, 970	16.277	25.768	44. 322	40. 901	34.882
	HT2RM2000984	38.635	39.635	42.628	14.734	10.729	39.002	24.661	39.000
	NT2RM2000994	38.405	43.907	36.416	29, 496	24.408	22. 384	18.679	31.517
	NT2RM2001004	74. 509	45.438	146. 622	36.919	35.918	125. 242	81.529	92.360
	NT2RM2001022	195. 677	346.056	350. 501	243.410	179.341	419.711	214.981	540.668
45	NT2RM2001035	23. 201	26.826	34. 867	15. 930	11.692	19.371	11.576	23.987
40	NT2RM2001038	18. 845	16.860	28. 577	14. 251	9.432	21.182	12.726	12.544
	NT2RH2001043		17. 293	22.001	11.462	11.232	18. 219	25.898	31, 106
		31.149		56.097	28. 126	32.674	61.600	49.621	79.938
	NT2RU2001050	101.638	45.617	<del></del>				35. 926	29.242
	NT2RM2001055	83.075	29.856	49. 927	15.739	32.251	60.461		14. 433
	MT2RM2001065	21.466	21.970	40. 152	20.006	27.398	26.370	15.034	
50	NT2RM2001075	366.658	258. 334	337.690	128. 945	166.931	370.161	257.064	228. 430
	NY 28112001083	230.683	79.913	107. 950	30. 576	63.142	203.365	79.590	24. 253
	MT 2RM2001100	182.772	114.627	137.289	65.878	54.062	141.899	155.507	119.434
	NT2RM2001105	101.949	70.116	95.624	50.863	39.812	104. 272	87.573	85. 122
	NT2RM2001109	48. 591	27.328	30.825	11.569	12.495	53.494	34.958	45. 222
	NT2RH2001110	99.871	68.967	152. 982	31.616	42.715	78.028	71.894	63.509
55	NT2RM2001125	57.602	33. 922	47.638	18.667	20.095	52.257	42.378	28. 204
	[11121112001120	31.002	1	1	1 .0.001	1			ii

Table 67

		en aca I	A4 F45 T	20 024 I	04 0C4 T	00 700 1	37 696 1	20 072	17.418
	NT2RM2001131	59.454	21.547	32.934	24.063	22.706	37.676	28.873	
j	NY2RH2001141	116.250	82.599	275.090	51.756	53.614	85.069	47. 274	83.199
	NT2RM2001152	20. 261	21.814	23. 297	10.506	9, 194	20.068	10.068	22.007
	NT2RH2001177	44.847	43. 449	52.307	26.604	19. 552	41.709	26.283	55. 231
	NT2RU2001194	164.727	54. 905	97. 293	28. 358	44. 057	146.597	99.019	118.606
	NT2RH2001195	36.939	36.245	34.818	15.750	15. 727	32.602	21.851	34. 274
	NT2RM2001196	125, 134	23.362	52, 729	15.781	26.090	77, 518	62.058	31.794
	NT2RM2001201	\$6.981	42.504	62.447	20. 139	31.351	68.607	32.835	44, 422
			70 746				40. 240	33.849	36. 497
	NT2RH2001221	65.764	32.746	40. 357	19.556	25. 529			
	NY2RM2001238	34. 807	25. 200	33. 023	13.254	14.872	43.011	20, 155	18. 493
	NT2RH2001243	50.316	49.076	42. 361	34. 148	33. 121	68.021	35.734	60.810
	NT2RM2001244	39.082	47.756	54.069	35. 242	30.728	59.908	22.778	50. 393
	NT2RM2001247	138. 825	184. 906	146, 564	65.082	57.954	94.133	78.544	136,745
	NT2RM2001256	28. 147	18.773	29. 336	14. 133	9. 881	8.739	16.106	25. 473
							11.919	14, 441	17.847
	NT2RH2001269	21.655	19.444	36.676	14. 235	17.978			
	HT2RM2001278	105. 133	67.683	225. 135	41.243	42.803	61.361	51.930	64. 103
	NT2RM2001291	21.264	19.798	31, 162	8.619	11.535	15. 945	16.243	12. 482
	NT2RM2001294	60.754	44.696	66. 102	25.820	20.715	42.950	28.321	33. 134
	NT2RM2001295	43.856	35. 189,	40.675	10.220	16.301	35.694	20.908	35. 879
	NT2RM2001302		16. 802	26.058	10. 228	12. 245	25. 513	14.404	12.416
		30.816		16 700					10. 145
	NY2R32001306	11.584	52. 176	16.722	6.379	6.616	13.560	8.347	
	HT2RM2001312	33.361	18.866	54. 572	11.148	10.119	13.848	8. 526	26.714
	NT2R12001319	13. 127	22.841	- 23.586	17.119	10. 492	18.998	4.495	36. 587
	NT2RM2001324	103.673	83.091	165, 198	32.861	22.836	56.112	31.793	39, 459
	NY2RM2001345	49. 634	25. 168	35.284	14.837	16. 900	100.618	25.540	19. 919
	NT2812001360	74. 152	33.097	38.122	17.360	16.021	50.562	31.265	21.915
						5. 734	26, 406	10,631	2. 394
	NT2RM2001370	28.821	12.859	21.986	6.327				
	四个2世2001391	16. 127	5.412	27.834	4.575	4. 553	14.188	3.910	9. 994
	MT2RM2001393	57. 930	25. 241	58.135	14.781	20. 544	47. 187	32.903	28. 104
	NT2RM2001420	17. 272	10.676	16.079	6.774	6.751	2.717	3.157	8. 464
	NT28112001423	17. 345	9.837	15. 261	12. 233	6. 527	15. 432	10.007	10. 935
	NT2RM2001424	196. 973	74. 966	136.019.	35. 222	48.814	142.268	95, 111	56, 187
			14. 500				227.572	99. 155	72. 372
	NT2RM2001482	265. 035	123.493	274. 926	59.811	62.022			
	NT2RM2001499	65. 942	48.790	62.383	28.605	19.730	68.321	23.722	26. 475
	NT2RM2001504	39. 282	24.742	30.958	9. 395	16. 991	46.880	13.034	16.709
	NT2RM2001524	24.755	14.244	24. 384	9.699	10.204	16.924	9.647	14. 539
	NY2RM2001530	5. 573	8.914	10.768	5.856	3, 286	9.623	4.337	7. 511
	NT2RM2001533	69. 137	57.026	127.055	29. 970	34.159	33.371	27.483	25. 268
	NT2822001540	65.400	54. 541	73.017	63. 277	35.636	49.097	31.308	76.346
							19.380	11.033	9. 485
	NT2RM2001544	18.067	19.624	25. 228	12.549	7.049			
	NT2RM2001547	22. 357	25.608	19.122	11.755	13.130	14.503	12.339	10.697
	NT2RY2001558	59.623	25.861	31.696	14. 111	16.568	53.758	34,606	18. 325
	NT2RM2001575	53. 128	46.425	111.368	27.392	24. 257	43.005	25. 405	24. 423
	NT2RM2001582	59.050	42.778	132.294	24.555	24.449	28. 347	22. 303	22. 397
	NT2R92001588	35. 342	21.815	27. 343	8.806	14. 132	21.498	16.451	22. 464
	NT2RM2001592	19.456	18. 542	28. 436	10.182	12.538	15.234	15. 478	15. 460
	NT2RM2001603			41.037	12.377	16.738	23. 117	21.517	12. 277
		42.456	15. 253		12 500	10.730		20.117	15. 459
	NT2RM2001605	60. 434	36, 233	43. 204	13.580	20.116	41. 260		
	NT2RM2001611	54.771	39.056	128. 984	17.180	24.100	40.047	19.191	16. 136
	NT2RM2001613	39. 500	22.894	27.579	12.321	11.577	26.696	21.149	24.773
	NY2RM2001626	202.358	40.774	93. 458	19.731	45.138	168.993	96.729	42.842
	NT2RM2001632	30. 160	45. 268	47. 586	25, 780	18.848	32,974	21, 939	45. 513
	NT2RH2001633		9. 885	12.546	7.571	6.017	11.226	7.294	20.798
		6.521					115.049	88. 246	58. 313
	NT2RM2001635	188.515	41.783	101.462	30.227	41.863			
	NT2RH2001638	26.880	23.087	31.788	15.679	14. 225	22.589	16.870	26. 264
	NT2RM2001637	13.020	5. 524	6.631	4.897	11.170	10,700	20.526	5. 331
	HY2RM2001639	71.531	28.740	32. 389	12.149	15.813	54.897	28.931	13. 443
	NT2RM2001841	39. 297	32, 462	49, 334	14.630	22,002	30.556	21.763	16.776
	101 2RU2001643	25. 535	12.621	15.764	6.658	12.027	21.274	22.136	12.847
				24. 507		6.636	18. 218	14.277	13.561
	NT28112001648	26.584	18.351						
	NT 2RM2001652	18.655	15.854	22. 304	6.782	9.644	25. 729	7.851	20. 144
	NT2RH2001659	16.893	10.861	16. 538	3.750	4.964	9. 228	6.172	11.278
	HT2RM2001660	17.414	13, 987	20.619	12.709	10,544	12.482	10.671	11.244
	NY2RH2001664	32.470	29, 186	27.804	16.171	15, 728	29.928	13.136	17.877
		1 -2. 4. 0	,			· · · · · · · · · · · · · · · · · · ·			

#### Table 68

HTTRENGOISES   83,125   51,356   52,221   12,055   14,144   18,196   46,704   15,988   HTRENGOISTO   35,448   70,552   70,752   71,777   75,452   67,725   75,514   50,752   71,777   75,452   77,775   75,527   71,777   75,452   77,775   75,527   71,777   75,452   77,775   75,527   77,777   75,527					I u	DIC OU			_	
		NT2RM2001668	89.325	61.356	52. 221	32.055	34. 144	88. 196		
		NT2RM2001670		20. 552	40. 552	17, 717	15. 452	67.725		
INTERNITORIES   5.784   7.472   11.234   3.584   7.484   5.095   5.298   21.547		NT2RM2001671	31, 368	15.752	21.018	19.630		62.746		
TYPENEZONISS   28,752   21,105   22,146   9,525   9,056   19,334   21,485   26,746   17,185   17,185   18,185   19,185   19,185   19,185   18,185   18,185   19,185   18,185		NT2RM2001675	7. 281	7.210	6.726	2.026	5.059	4.678		
INTEREDUCTIONS   32, 313   25, 779   43, 734   11, 155   10, 491   20, 228   33, 991     INTEREDUCTIONS   82, 268   103, 407   49, 403   48, 305   60, 147   82, 201   35, 207   61, 706     INTEREDUCTIONS   146, 791   45, 033   31, 099   28, 885   31, 134   11, 185   65, 047   78, 79, 79, 79, 79, 79, 79, 79, 79, 79, 79		NT2RM2001681	5.784	7.472	11. 234	3.584	7.454	5.095		
STRENGOLISES   32, 233   25, 279   43, 734   11, 154   11, 156   30, 491   20, 238   33, 91     STRENGOLISES   62, 086   103, 450   329, 341   48, 305   60, 347   85, 201   35, 503   11, 706     STRENGOLISES   101, 355   65, 027   88, 822   31, 652   35, 701   110, 799   51, 158   32, 399     STRENGOLISES   24, 377   20, 994   25, 919   13, 654   894   19, 112   14, 447   42, 307     STRENGOLISES   24, 377   20, 994   25, 919   13, 654   89, 94   19, 112   14, 447   42, 307     STRENGOLISES   24, 377   20, 994   25, 919   13, 654   89, 94   19, 112   14, 447   42, 307     STRENGOLISES   75, 478   27, 887   50, 059   14, 943   24, 128   36, 190   27, 006   62, 522     STRENGOLITOR   50, 1932   27, 867   50, 059   14, 943   24, 128   36, 190   27, 006   62, 522     STRENGOLITOR   57, 478   57, 678   57, 778   57, 273   30, 379   105, 548   62, 864   26, 547     STRENGOLITOR   105, 052   48, 161   57, 895   15, 177   34, 379   105, 548   62, 864   26, 050     STRENGOLITOR   27, 206   27, 858   28, 558   16, 545   51, 327   33, 590   48, 519     STRENGOLITOR   27, 206   27, 859   28, 859   16, 542   59, 397   19, 551   31, 397   38, 699   88, 079   89, 079   8		NY2RM2001685	28.752	21.105	22. 145	9. 525	9.058	19. 334		
STRENGOLISS   02.088   103.403   239.543   46.305   60.347   69.201   35.903   81.705     RTRENGOLISS   101.355   65.027   68.822   31.652   35.701   101.799   51.358   52.359     RTRENGOLISS   446.791   45.033   83.099   28.886   31.134   111.891   66.042   79.212     RTRENGOLISS   446.791   45.033   83.099   28.886   31.134   111.891   66.042   79.212     RTRENGOLISS   446.791   45.033   83.099   28.886   31.134   111.891   66.042   79.212     RTRENGOLISS   55.393   72.887   50.059   14.943   42.172   36.894   19.12   14.447   42.307     RTRENGOLISS   55.393   72.887   50.059   14.943   42.172   48.190   27.006   62.522     RTRENGOLISS   75.276   62.308   444.702   48.187   41.966   55.679   28.916   57.741     RTRENGOLITIS   12.4088   99.615   122.970   48.569   68.313   188.154   109.177   48.112     RTRENGOLITIS   28.058   99.615   122.970   48.569   68.313   188.154   109.177   48.112     RTRENGOLITIS   28.058   99.615   122.970   48.569   68.313   188.154   109.177   48.112     RTRENGOLITIS   27.044   41.046   48.272   22.655   68.313   88.154   109.177   48.112     RTRENGOLITIS   77.064   41.046   48.272   22.655   6.545   53.455   13.27   33.590   8.807   9.491     RTRENGOLITIS   77.064   41.046   48.272   22.655   6.545   13.655   13.27   33.593   8.597   13.857   12.425     RTRENGOLITIS   31.218   31.218   31.518   31.518   31.218   31.			35, 233	25, 279	43.734	11. 154	11.656	30. 491		
RTZREZGOTEGE   OLI 355   65, OZ7   68, 822   31, 652   35, TOL   110, 799   51, 358   32, 798   RTZREZGOTEGE   46, 791   46, 793   47, 793   42, 999   28, 886   31, 134   11, 891   66, OZ4   79, 924   79, 793   79,		NT2RM2001695	82.068		239. 543	46. 305	60.347	69.201	35. 903	
NTZERIZOOTISSB   446,79  45,033   83,099   28,866   31,134   11,891   66,042   79,232     RTZERIZOOTISSB   74,377   03,944   25,994   25,994   25,994   25,994   25,994   25,994   27,813   27,813   24,85   43,74   42,307     RTZERIZOOTIOS   17,314   8,383   12,975   3,702   1,312   7,813   9,485   4,374     RTZERIZOOTIOS   55,476   62,306   144,022   48,167   41,966   55,679   28,916   57,741     RTZERIZOOTIOS   75,476   62,306   144,022   48,167   41,966   55,679   28,916   57,741     RTZERIZOOTIOS   73,676   73,916   73,550   73,450   73,451   73,415					68. 822	31.652	35.701	110.799	51.358	
RTZERZGOTIGES   24,377   20,994   25,919   13,654   8,984   9,132   14,447   42,307   RTZERZGOTIGES   14,714   50,393   27,856   50,059   14,943   24,129   36,190   27,006   62,526   RTZERZGOTIGES   75,676   27,857   50,059   14,943   24,129   36,190   27,006   62,527   RTZERZGOTIGES   75,676   27,857   50,059   14,943   24,129   36,190   27,006   62,576   77,741   77,741   77,757   RTZERZGOTIGES   75,676   75,976   75,767   75,767   75,767   75,767   75,777   74,777   74,117   75,777   75,7		NT2RM2001698	146.791	45.033	83.099	28.886	31.134	111.891	66.042	
NTTERIZEDOTOTOD			24, 737	20. 994	25. 919	13.654	8.984	19.132		
### ### ### ### ### ### ### ### ### ##			14.734	8. 383	12.975	3.702	1.312	7.813		
		NT2RH2001704	50. 393	27.867	50.059	14, 943				
ST72812001716   294, 058   93, 615   122, 970   48, 559   68, 313   188, 154   109, 177   48, 112     ST72812001718   20, 1352   14, 923   16, 575   7, 213   8, 937   39, 809   8, 807   9, 491     ST2812001723   20, 1352   14, 923   16, 575   7, 213   8, 937   39, 809   8, 807   9, 491     ST2812001730   27, 206   62, 229   24, 855   8, 552   9, 397   19, 553   31, 897   12, 427     ST2812001733   27, 206   62, 229   18, 054   7, 967   10, 452   22, 532   14, 233   28, 610     ST2812001734   31, 219   15, 575   27, 495   8, 999   13, 556   19, 966   11, 234   20, 203     ST2812001735   102, 208   95, 533   95, 880   48, 800   50, 926   85, 016   46, 946   86, 945     ST2812001735   102, 208   95, 533   95, 880   48, 800   50, 926   85, 016   46, 946   86, 945     ST2812001755   102, 208   95, 533   95, 880   48, 800   50, 926   85, 016   46, 946   86, 945     ST2812001767   31, 834   16, 852   22, 405   14, 516   36, 220   11, 083   21, 129     ST2812001767   30, 834   31, 815   59, 888   20, 85, 959   13, 614   48, 508   313, 42   21, 29     ST28120017761   33, 843   31, 815   59, 888   20, 959   91, 855   34, 474   48, 508   37, 342   20, 583     ST2812001768   14, 334   16, 852   22, 405   54, 516   36, 420   10, 682   25, 114   20, 583     ST2812001769   36, 852   28, 592   43, 280   11, 529   16, 235   41, 973   21, 095   36, 897     ST2812001767   33, 843   31, 815   59, 888   20, 989   91, 855   34, 937   21, 209   36, 897     ST2812001768   14, 334   16, 852   22, 405   14, 516   7, 327   11, 583   21, 129     ST28120017791   33, 843   31, 815   59, 888   20, 989   91, 855   34, 937   27, 736     ST2812001792   49, 540   17, 667   39, 944   11, 809   19, 235   60, 433   38, 300   42, 078     ST2812001792   73, 444   44, 32, 799   54, 722   44, 888   28, 332   74, 431   52, 678   40, 155     ST28122001792   73, 744   44, 52, 799   54, 722   44, 888   28, 332   74, 431   72, 678   64, 528     ST28122001893   30, 30, 34   36, 30   39, 942   31, 894   31, 895   32, 30   30, 30   32, 30   32, 30   32, 30			75.476		144. 702	48. 167	41.996	55. 679		
STZERZDO1718		NT2RH2001714	14.876	12.916	25.654	7.345	10.946			
NTZRIZZO17737   20. 352		HT2RH2001716	294.058	99.615	122.970					
NTZRIZZO01737   57,044   41,046   46,272   22,685   16,545   51,332   33,590   46,539   8172RIZZO01738   25,035   6,229   818,054   7,367   10,452   22,532   14,238   26,610   8172RIZZO01748   25,035   6,229   818,054   7,367   10,452   22,532   14,238   26,610   8172RIZZO01743   41,699   57,379   68,633   29,155   63,474   48,600   37,342   50,583   8172RIZZO01755   102,308   95,533   95,880   48,800   50,926   85,016   46,946   58,535   8172RIZZO01765   7,310   22,525   20,809   5,477   6,161   36,420   11,083   21,129   8172RIZZO01765   7,310   22,525   20,809   5,477   6,161   36,420   11,083   21,129   8172RIZZO01765   17,310   22,525   20,809   5,477   6,161   36,420   11,083   21,129   8172RIZZO01768   41,334   16,852   22,405   14,516   7,327   13,653   4,371   27,736   8172RIZZO01778   4,653   9,177   12,741   9,99   6,577   9,552   26,114   70,587   8172RIZZO01778   4,433   9,177   12,741   9,99   6,577   9,552   26,613   4,371   27,736   8172RIZZO01778   4,434   32,799   54,772   4,488   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,799   54,772   4,888   28,332   7,443   52,678   40,155   8172RIZZO01795   73,444   32,793   73,878   22,935   44,781   108,971   66,672   68,900   8172RIZZO01795   73,748   74,748		NT2RM2001718	109.052	48.161		15, 717				
NYZRIZ201730		NT2RM2001723	20.352		16.575		8. 937			
NTZRIZ2001738   25.036   6.229   18.054   7.967   10.452   22.532   14.238   26.610   NTZRIZ2001743   31.219   15.575   27.495   8.999   13.856   13.956   27.123   40.203   NTZRIZ2001753   41.699   57.379   56.833   29.155   36.474   48.608   37.342   50.583   NTZRIZ2001755   102.308   95.543   95.880   48.800   50.926   85.016   46.946   58.515   NTZRIZ2001765   17.310   22.525   23.809   15.276   61.235   41.973   21.095   36.877   17.276   22.525   23.809   5.472   5.161   36.420   11.083   21.129   HTZRIZ2001765   17.310   22.525   23.809   5.472   5.161   36.420   11.083   21.129   HTZRIZ2001765   14.334   16.852   22.405   14.516   7.327   13.653   4.371   27.736   HTZRIZ2001776   14.334   16.852   22.405   14.516   7.327   13.653   4.371   27.736   HTZRIZ2001771   31.863   31.815   59.888   20.939   19.261   40.662   26.114   70.587   HTZRIZ2001777   14.653   9.177   12.741   0.999   6.577   9.552   8.651   6.255   8.5516   8.728   8.7		NT2RM2001727		41.046						
NTZRIZO01743   31.219   15.575   27.495   8.999   13.856   19.966   21.123   40.203     NTZRIZO01755   102.308   95.543   95.880   48.800   50.926   85.016   46.946   58.535     NTZRIZO01760   36.852   29.592   43.280   11.529   16.235   41.973   21.095   36.887     NTZRIZO01765   103.038   95.543   95.880   48.800   50.926   85.016   46.946   58.535     NTZRIZO01765   103.038   95.864   47.800   11.529   16.235   41.973   21.095   36.887     NTZRIZO01765   103.038   98.824   244.752   82.225   86.662   313.630   261.579   156.449     NTZRIZO01765   14.334   16.852   22.405   14.516   7.327   13.653   4.371   77.746     NTZRIZO01777   33.884   31.815   55.888   20.939   19.261   40.682   25.114   70.587     NTZRIZO01777   33.884   31.815   55.888   20.939   19.261   40.682   25.114   70.587     NTZRIZO01774   49.540   17.667   39.944   11.809   19.255   60.433     NTZRIZO01794   49.540   17.667   39.944   11.809   19.255   60.433     NTZRIZO01794   49.540   17.667   39.944   11.809   19.255   60.433     NTZRIZO01795   73.444   32.799   54.722   41.888   28.32   74.43   12.678   40.155     NTZRIZO01795   73.444   32.799   54.722   41.888   28.32   74.43   12.678   40.155     NTZRIZO01795   73.744   32.799   54.722   41.888   28.32   74.43   12.678   40.155     NTZRIZO01797   77.70   23.911   46.302   31.918   15.965   38.330   15.267   61.400     NTZRIZO01797   77.770   23.911   46.302   31.918   15.965   38.330   15.267   61.400     NTZRIZO01800   30.838   39.806   27.862   15.915   15.950   33.330   15.267   61.400     NTZRIZO01805   10.973   6.105   2.1287   5.928   6.677   77.463   7.004   10.764     NTZRIZO01805   10.973   6.105   2.1287   5.928   6.677   77.463   7.004   10.764     NTZRIZO01805   10.973   6.105   2.2867   3.095   5.188   10.179   14.993   12.571   5.587     NTZRIZO01808   37.838   39.806   27.862   3.955   5.468   10.179   14.993   12.571   5.587     NTZRIZO01808   37.838   39.806   27.862   3.959   5.188   10.179   10.847   37.948   37.948   37.948   37.948   37.948   37.948   37.948										
NTTRIEDUTITS										
STERNIZUOT755   102. 308   35. 543   35. 880   48. 800   50. 926   85. 016   46. 946   58. 515										
NT2RE2001765   17. 310   22. 525   20. 809   5. 472   6. 215   36. 420   11. 083   21. 123     NT2RE2001767   17. 310   22. 525   20. 809   5. 472   6. 161   36. 420   11. 083   21. 123     NT2RE2001768   14. 314   16. 852   22. 405   14. 516   7. 327   13. 653   4. 371   27. 736     NT2RE2001771   33. 884   31. 815   53. 888   20. 959   91. 261   40. 662   26. 114   70. 581     NT2RE2001778   14. 653   3. 177   12. 741   0. 999   6. 577   9. 552   8. 651   6. 225     NT2RE2001778   49. 540   17. 667   39. 944   11. 809   19. 235   60. 433   38. 302   42. 076     NT2RE2001784   31. 529   23. 807   34. 905   9. 620   16. 512   6. 574   41. 749   17. 088     NT2RE2001785   73. 444   32. 799   54. 722   14. 868   28. 332   74. 431   52. 678   40. 155     NT2RE2001792   82. 550   48. 689   54. 661   13. 880   26. 470   67. 309   56. 934   51. 170     NT2RE2001793   130. 534   65. 803   79. 887   22. 935   40. 781   108. 971   66. 672   68. 900     NT2RE2001090   32. 076   15. 750   32. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001806   10. 973   6. 105   22. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001805   10. 973   6. 105   22. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001805   10. 973   6. 105   22. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001805   10. 973   6. 105   22. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001805   10. 973   6. 105   22. 039   9. 323   10. 196   25. 569   24. 848   32. 579     NT2RE2001805   10. 973   6. 105   52. 678   52. 678   10. 173   17. 178   25. 827     NT2RE2001805   10. 973   6. 105   52. 679   5. 168   10. 179   17. 178   25. 827     NT2RE2001805   10. 973   6. 105   52. 678   52. 678   13. 194   12. 194										
NT2R12001765										
NTZERIZUDIY67   S07. 383   198. 624   244.752   82. 225   86. 662   313. 630   261. 679   156. 449     NTZERIZUDIY68   14. 334   16. 852   22. 405   14. 516   7. 327   13. 653   4. 371   27. 736     NTZERIZUDIY71   34. 653   9. 177   12. 741   0. 999   6. 577   9. 552   8. 651   6. 525     NTZERIZUDIY62   49. 540   17. 667   39. 944   11. 809   19. 225   60. 433   38. 002   42. 078     NTZERIZUDIY64   31. 529   23. 807   34. 905   9. 620   16. 512   26. 777   41. 749   17. 008     NTZERIZUDIY65   73. 444   32. 799   54. 722   14. 868   28. 332   74. 431   52. 678   40. 155     NTZERIZUDIY65   73. 444   32. 799   54. 762   13. 880   26. 470   67. 300   56. 934   51. 170     NTZERIZUDIY65   82. 550   48. 689   54. 661   13. 880   26. 470   67. 300   56. 934   51. 170     NTZERIZUDIY67   17. 770   23. 911   46. 302   31. 918   15. 955   38. 330   15. 267   61. 440     NTZERIZUDIORO   32. 076   15. 750   32. 039   9. 323   10. 196   25. 569   24. 488   32. 579     NTZERIZUDI803   18. 883   19. 805   27. 862   15. 915   15. 790   17. 317   12. 178   25. 827     NTZERIZUDI805   10. 973   6. 105   12. 362   3. 395   7. 748   17. 42   7. 464   10. 576     NTZERIZUDI805   13. 11. 155   10. 752   12. 187   5. 926   6. 671   17. 463   7. 004   10. 764     NTZERIZUDI813   11. 155   10. 752   12. 187   5. 926   6. 671   17. 463   7. 004   10. 764     NTZERIZUDI813   13. 814   13. 268   12. 712   4. 452   7. 791   10. 847   8. 727   7. 819     NTZERIZUDI815   27. 524   37. 936   22. 505   15. 145   17. 486   21. 050   17. 161   33. 945     NTZERIZUDI825   27. 524   37. 936   22. 505   15. 145   17. 486   21. 050   17. 161   33. 945     NTZERIZUDI826   33. 814   13. 268   12. 712   4. 452   7. 791   10. 847   8. 727   7. 819     NTZERIZUDI815   52. 202   39. 752   63. 088   24. 308   18. 778   32. 221   26. 626   85. 665     NTZERIZUDI836   33. 814   13. 268   12. 712   4. 452   7. 791   10. 847   8. 727   7. 819     NTZERIZUDI837   53. 30. 60   62. 176   53. 785   63. 088   24. 308   18. 778   32. 221   17. 162   33. 355										
NT2REP2001768										
NTZRIZO01777										
NTZRIZEDOTYPE   44, 653   3, 177   12, 741   0, 999   6, 577   9, 552   8, 651   6, 525										
NTZREJZ001782										
NTZRHZ001792   31.529   23.807   34.905   9.520   16.512   26.774   14.749   17.008   NTZRHZ001795   73.444   32.799   54.722   14.868   28.332   74.431   52.678   40.155   NTZRHZ001795   87.544   35.279   54.722   14.868   28.332   74.431   52.678   40.155   NTZRHZ001795   130.534   65.803   79.887   22.935   40.781   108.971   66.672   68.900   NTZRHZ001797   17.770   23.911   46.302   31.918   15.965   38.330   15.267   61.440   NTZRHZ001800   32.076   15.750   32.039   9.323   10.196   25.569   24.848   32.579   NTZRHZ001803   18.883   19.806   27.862   15.915   15.790   17.317   12.178   25.827   NTZRHZ001805   10.973   6.105   12.362   3.395   7.748   17.242   7.464   10.576   NTZRHZ001805   41.604   28.683   30.345   12.360   14.554   35.269   18.192   22.416   NTZRHZ001813   11.155   10.752   12.187   5.926   6.671   17.463   7.004   10.764   NTZRHZ001818   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747   NTZRHZ001818   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747   NTZRHZ001825   27.524   37.936   22.505   15.145   17.486   21.050   17.161   33.945   NTZRHZ001825   27.524   37.936   22.505   15.145   17.486   21.050   17.161   33.945   NTZRHZ001825   27.524   37.936   22.505   15.145   17.486   21.050   17.161   33.945   NTZRHZ001835   53.715   31.908   39.273   13.944   21.144   27.291   25.952   18.816   NTZRHZ001855   53.006   24.176   29.953   16.912   19.394   23.562   31.355   25.910   NTZRHZ001855   33.006   24.176   29.953   16.912   19.394   23.562   31.355   25.910   NTZRHZ001867   30.838   22.957   35.457   44.948   16.183   32.799   17.562   46.800   NTZRHZ001865   33.026   24.176   29.953   16.912   19.394   23.562   31.355   25.910   NTZRHZ001865   33.026   24.176   29.953   16.912   19.394   23.562   31.355   25.910   NTZRHZ001867   30.838   22.957   35.457   44.948   64.737   231.277   44.776   47.729   NTZRHZ001867   30.838   22.957   35.457   44.948   64.737   231.277   45.766   46.800   NTZRHZ001866   31.621   99.975   63.088   24.308										
NTZREZ2001792										
NTZREZOD1792   82.550   48.689   54.661   13.880   26.470   67.309   56.934   51.170     NTZREZOD1795   130.534   65.803   79.887   22.935   40.781   108.971   66.672   68.900     NTZREZOD1797   17.770   23.911   46.302   31.918   15.965   38.330   15.267   61.440     NTZREZOD1800   32.076   15.750   32.039   9.323   10.196   25.569   24.848   32.579     NTZREZOD1803   18.883   19.806   27.862   15.915   15.790   17.317   12.178   25.827     NTZREZOD1805   10.973   6.105   12.362   3.395   7.748   17.242   7.464   10.576     NTZREZOD1806   41.604   28.683   30.345   12.360   14.554   35.269   18.192   22.416     NTZREZOD1806   41.604   28.683   30.345   12.360   14.554   35.269   18.192   22.416     NTZREZOD1813   11.155   10.752   12.187   5.926   6.671   17.463   7.004   10.764     NTZREZOD1814   16.422   18.276   19.059   5.168   10.179   14.993   12.571   9.506     NTZREZOD1818   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747     NTZREZOD1825   27.524   37.936   22.505   15.145   17.486   21.050   17.161   33.945     NTZREZOD1825   27.524   37.936   22.505   15.145   17.486   21.050   17.161   33.945     NTZREZOD1830   53.715   31.908   39.273   13.944   12.144   27.291   25.952   18.816     NTZREZOD1830   53.715   31.908   39.273   13.944   12.144   27.291   25.952   18.816     NTZREZOD1855   33.026   24.176   29.953   16.912   19.394   23.526   31.355   25.910     NTZREZOD1855   33.026   24.176   29.953   16.912   19.394   23.526   31.355   25.910     NTZREZOD1855   33.026   24.176   29.953   16.912   19.394   23.526   31.355   25.910     NTZREZOD1858   33.026   24.176   29.953   16.912   19.394   23.526   31.355   25.910     NTZREZOD1859   129.599   162.083   180.222   173.694   64.737   231.277   145.176   147.129     NTZREZOD1869   129.599   162.083   180.222   173.694   64.737   231.277   145.176   147.129     NTZREZOD1869   129.599   162.083   180.222   173.694   64.737   231.277   145.176   147.129     NTZREZOD1868   31.621   19.917   31.650   19.861   14.683   19.395   24.61										
NTZRHIZODIT9S   130.534   65.803   79.887   22.935   40.781   108.971   66.672   68.900     NTZRHIZODIT9S   17.770   23.911   46.302   31.918   15.965   38.330   15.267   61.440     NTZRHIZODIBOO   32.076   15.750   32.039   9.323   10.196   25.569   24.848   32.579     NTZRHIZODIBOO   318.883   19.806   27.862   15.915   15.790   17.317   12.178   25.827     NTZRHIZODIBOS   10.973   6.105   12.362   3.395   7.748   17.242   7.464   10.576     NTZRHIZODIBOS   41.604   28.683   30.345   12.360   14.554   35.269   18.192   22.416     NTZRHIZODIBIA   11.155   10.752   12.187   5.926   6.671   17.463   7.004   10.764     NTZRHIZODIBIA   11.155   10.752   12.187   5.926   6.671   17.463   7.004   10.764     NTZRHIZODIBIB   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747     NTZRHIZODIBIB   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747     NTZRHIZODIBIB   37.340   15.047   25.378   7.050   13.614   28.082   23.903   16.747     NTZRHIZODIBIZ   38.14   13.268   12.712   4.562   7.791   10.847   8.727   7.819     NTZRHIZODIBIZ   68.657   29.677   30.202   9.749   22.522   37.241   30.727   18.205     NTZRHIZODIBIZ   68.657   29.677   30.202   9.749   22.522   37.241   30.727   18.205     NTZRHIZODIBIS   53.715   31.908   39.273   33.944   12.144   27.291   25.952   18.816     NTZRHIZODIBIS   52.202   39.752   63.088   24.308   18.778   32.821   26.626   85.665     NTZRHIZODIBIS   52.202   39.752   63.088   24.308   18.778   32.821   26.626   85.665     NTZRHIZODIBIS   52.202   39.752   63.088   24.308   18.778   32.821   26.626   85.665     NTZRHIZODIBIS   14.477   14.016   20.104   6.241   7.997   18.463   6.634   17.934     NTZRHIZODIBIS   30.332   1475   45.925   173.694   64.737   231.277   145.176   147.129     NTZRHIZODIBIS   52.202   39.752   63.088   24.308   18.778   32.277   145.176   147.129     NTZRHIZODIBIS   52.506   54.176   29.953   16.912   19.394   23.662   31.521   16.626   85.665     NTZRHIZODIBIS   52.506   54.649   14.914   59.041   11.657   28.809   14										
NTZRHZQO1190										
NTZRH2001800   32.076   15.750   32.039   9.323   10.196   25.569   24.848   32.579										61.440
RTZRE2001803   18.883   19.806   27.862   15.915   15.790   17.317   12.178   25.827	•									32.579
NTZRIZ001805   10.973   6.105   12.362   3.395   7.748   17.242   7.464   10.576								17.317	12.178	25. 827
NTZRH2001836								17.242	7. 464	
NTZRH2001813						12, 360	14.554	35. 269	18. 192	22.416
RTZRHZ001818   37,340   15,047   25,378   7,050   13,614   28,082   23,903   16,747     RTZRHZ001823   13,814   13,268   12,712   4,562   7,791   10,847   8,727   7,819     RTZRHZ001825   27,524   37,936   22,505   15,145   17,486   21,050   17,161   33,945     RTZRHZ001832   68,657   29,677   30,202   9,749   22,522   37,241   30,727   18,205     RTZRHZ001839   53,715   31,908   39,273   13,944   12,144   27,291   25,952   18,816     RTZRHZ001840   108,411   98,429   259,021   48,048   32,857   58,314   28,523   37,338     RYZRHZ001851   52,202   39,752   63,088   24,308   18,778   32,821   26,626   85,666     RTZRHZ001855   33,026   24,176   29,953   16,912   19,394   23,562   31,355   25,910     RTZRHZ001867   30,838   22,957   35,457   14,948   16,183   32,799   17,562   46,800     RTZRHZ001867   30,838   22,957   35,457   14,948   16,183   32,799   17,562   46,800     RTZRHZ001869   129,599   162,083   180,222   173,694   64,737   231,277   145,176   147,129     RTZRHZ001869   31,621   19,917   31,650   19,861   14,683   19,396   24,619   18,912     RTZRHZ001886   31,621   19,917   31,650   19,861   14,683   19,396   24,619   18,912     RTZRHZ001886   5201,332   1475,462   2605,875   738,729   3013,651   5911,225   5347,627   1306,593     RTZRHZ001896   5201,332   1475,462   2605,875   738,729   3013,651   5911,225   5347,627   1306,593     RTZRHZ001896   5201,332   1475,462   2605,875   738,729   3013,651   5911,225   5347,627   1306,593     RTZRHZ001896   5201,332   1475,462   2605,875   738,729   3013,651   5911,225   5347,627   1306,593     RTZRHZ001936   53,243   40,127   55,162   28,793   22,732   77,356   28,595   48,438     RTZRHZ001935   36,519   23,148   18,415   4,134   11,165   15,562   19,141   10,042     RTZRHZ001935   36,519   23,148   18,415   4,134   11,165   15,562   19,141   10,042     RTZRHZ001935   36,519   23,148   18,415   4,134   11,165   15,562   19,141   10,042     RTZRHZ001935   23,961   5,651   21,192   6,301   5,377   17,197   7,025   5,482						5.926	6.671	17.463		
NT2RM2001823 13. 814 13. 268 12.712 4.562 7.791 10.847 8.727 7.819 NT2RM2001825 27. 524 37. 936 22.505 15. 145 17. 486 21.050 17. 161 33. 945 NT2RM2001832 68. 657 29. 677 30. 202 9. 749 22. 522 37. 241 30. 727 18. 205 NT2RM2001839 53. 715 31. 908 39. 273 13. 944 12. 144 27. 291 25. 952 18. 816 NT2RM2001840 108. 411 98. 429 259. 021 48. 048 32. 857 58. 314 28. 523 37. 338 NY2RM2001855 52. 202 39. 752 63. 088 24. 308 18. 778 32. 821 26. 626 85. 666 NT2RM2001855 33. 026 24. 176 29. 953 16. 912 19. 394 23. 562 31. 355 25. 910 NT2RM2001867 30. 838 22. 957 35. 457 14. 948 16. 183 32. 799 17. 562 46. 800 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 739 71 18. 463 6. 634 17. 934 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 739 18. 463 6. 634 17. 934 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 739 18. 463 6. 634 17. 934 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 739 18. 463 6. 634 17. 934 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001869 31. 621 19. 917 31. 650 19. 861 14. 683 19. 396 24. 619 18. 912 NT2RM2001869 5201. 332 1475. 462 2605. 875 738. 729 3013. 651 5911. 225 5347. 627 1306. 593 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 239 7. 418 7. 583 3. 383 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001935 36. 519 23. 148 18. 415 4. 134 11. 165 15. 562 19. 141 10. 042 NT2RM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NT2RM2001938 23. 961 5. 651 21. 192 6. 301 5. 377 17. 197 7. 025 5. 482		NT2RM2001814	16.422	18. 276	19.059	5. 168				
NT2RM2001825 27. 524 37. 936 22. 505 15. 145 17. 486 21.050 17. 161 33. 945 NT2RM2001832 68. 657 29. 677 30. 202 9. 749 22. 522 37. 241 30. 727 18. 205 NT2RM2001839 53. 715 31. 908 39. 273 13. 944 12. 144 27. 291 25. 952 18. 816 NT2RM2001840 108. 411 98. 429 259. 021 48. 048 32. 857 58. 314 28. 523 37. 338 NY2RM2001855 52. 202 39. 752 63. 088 24. 308 18. 778 32. 821 26. 626 85. 666 NT2RM2001855 33. 026 24. 176 29. 953 16. 912 19. 394 23. 562 31. 355 25. 910 NT2RM2001867 30. 838 22. 957 35. 457 14. 948 16. 183 32. 799 17. 562 46. 800 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001895 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001893 42. 649 14. 914 59. 041 11. 657 28. 809 14. 670 17. 172 5. 336 NT2RM2001895 31. 621 19. 917 31. 650 19. 861 14. 683 19. 396 24. 619 18. 912 NT2RM2001895 620. 332 1475. 462 2605. 875 738. 729 3013. 651 5911. 225 5347. 627 1306. 593 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 838 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001935 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NT2RM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NT2RM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NT2RM2001938 23. 961 5. 651 21. 192 6. 301 5. 377 17. 197 7. 0.25 5. 482				15. 047						
NTZRM2001832 68.657 29.677 30.202 9.749 22.522 37.241 30.727 18.205 NTZRM2001839 53.715 31.908 39.273 13.944 12.144 27.291 25.952 18.816 NTZRM2001840 108.411 98.429 259.021 48.048 32.857 58.314 28.523 37.338 NTZRM2001851 52.202 39.752 63.088 24.308 18.778 32.821 26.626 85.666 NTZRM2001853 33.026 24.176 29.953 16.912 19.394 23.562 31.355 25.910 NTZRM2001867 30.838 22.957 35.457 14.948 16.183 32.799 17.562 46.800 NTZRM2001869 129.599 162.083 180.222 173.694 64.737 231.277 145.176 147.129 NTZRM2001879 14.477 14.016 20.104 6.241 7.997 18.463 6.634 17.934 NTZRM2001888 31.621 19.917 31.650 19.861 14.683 19.396 24.619 18.912 NTZRM2001885 31.621 19.917 31.650 19.861 14.683 19.396 24.619 18.912 NTZRM2001885 5201.332 1475.462 2605.875 738.729 3013.651 5911.225 5347.627 1306.593 NTZRM2001902 9.512 5.176 9.030 3.230 3.539 7.418 7.583 3.383 NTZRM2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438 NTZRM2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 NTZRM2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NTZRM2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NTZRM2001938 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NTZRM2001938 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NTZRM2001938 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419										
NTZRM2001839   53, 715   31, 908   39, 273   13, 944   12, 144   27, 291   25, 952   18, 816   NTZRM2001840   108, 411   98, 429   259, 021   48, 048   32, 857   58, 314   28, 523   37, 338   NTZRM2001851   52, 202   39, 752   63, 088   24, 308   18, 778   32, 821   26, 626   85, 665   NTZRM2001855   33, 026   24, 176   29, 953   16, 912   19, 394   23, 562   31, 355   25, 910   NTZRM2001865   30, 838   22, 957   35, 457   14, 948   16, 183   32, 799   17, 562   46, 800   NTZRM2001869   129, 599   162, 083   180, 222   173, 694   64, 737   231, 277   145, 176   147, 129   NTZRM2001879   14, 477   14, 016   20, 104   6, 241   7, 997   18, 463   6, 634   17, 934   NTZRM2001803   42, 649   14, 914   59, 041   11, 657   28, 809   14, 670   17, 172   5, 396   NTZRM2001806   31, 621   19, 917   31, 650   19, 861   14, 683   19, 396   24, 619   18, 912   NTZRM2001896   31, 621   19, 917   31, 650   19, 861   14, 683   19, 396   24, 619   18, 912   NTZRM2001896   5201, 332   1475, 462   2605, 875   738, 729   3013, 651   5911, 225   5347, 627   1306, 593   NTZRM2001902   9, 512   5, 176   9, 030   3, 230   3, 539   7, 418   7, 583   3, 383   NTZRM2001902   9, 512   5, 176   9, 030   3, 230   3, 539   7, 418   7, 583   3, 383   NTZRM2001903   63, 243   40, 127   55, 162   28, 793   22, 732   77, 356   28, 595   48, 438   NTZRM2001935   63, 243   40, 127   55, 162   28, 793   22, 732   77, 356   28, 595   48, 438   NTZRM2001935   63, 519   23, 148   18, 415   4, 134   11, 165   15, 562   19, 141   10, 042   NTZRM2001935   78, 536   47, 939   51, 879   18, 980   21, 013   48, 576   35, 554   52, 419   NTZRM2001935   78, 536   47, 939   51, 879   18, 980   21, 013   48, 576   35, 554   52, 419   NTZRM2001935   78, 536   47, 939   51, 879   18, 980   21, 013   48, 576   35, 554   52, 419   NTZRM2001935   78, 536   47, 939   51, 879   18, 980   21, 013   48, 576   35, 554   52, 419   NTZRM2001935   78, 536   47, 939   51, 879   18, 980   21, 013   48, 576   35, 554   52, 419   NTZRM2001935   78, 536   47, 939   51, 879   1		***************************************								
NT2RM2001840 108. 411 98. 429 259.021 48. 048 32. 857 58. 314 28. 523 37. 338 NY2RM2001851 52. 202 39. 752 63. 088 24. 308 18. 778 32. 821 26. 626 85. 666 NT2RM2001855 33. 026 24. 176 29. 953 16. 912 19. 394 23. 562 31. 355 25. 910 NT2RM2001867 30. 838 22. 957 35. 457 14. 948 16. 183 32. 799 17. 562 46. 800 NT2RM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NT2RM2001879 14. 477 14. 016 20. 104 6. 241 7. 997 18. 463 6. 634 17. 934 NT2RM2001803 42. 649 14. 914 59. 041 11. 657 28. 809 14. 670 17. 172 5. 396 NT2RM2001886 31. 621 19. 917 31. 650 19. 861 14. 683 19. 396 24. 619 18. 912 NT2RM2001886 31. 621 19. 917 31. 650 19. 861 14. 683 19. 396 24. 619 18. 912 NT2RM2001886 5201. 332 1475. 462 2605. 875 738. 729 3013. 651 5911. 225 5347. 627 1306. 593 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NT2RM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 385 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RM2001935 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 418 NT2RM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 418 NT2RM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 418 NT2RM2001938 23. 961 5. 651 21. 192 6. 301 5. 377 17. 197 7. 0.25 5. 482										
NYZRM2001851 52. 202 39. 752 63. 088 24. 308 18. 778 32. 821 26. 626 85. 666 NYZRM2001855 33. 026 24. 176 29. 953 16. 912 19. 394 23. 562 31. 355 25. 910 NYZRM2001867 30. 838 22. 957 35. 457 14. 948 16. 183 32. 799 17. 562 46. 800 NYZRM2001869 129. 599 162. 083 180. 222 173. 694 64. 737 231. 277 145. 176 147. 129 NYZRM2001879 14. 477 14. 016 20. 104 6. 241 7. 997 18. 463 6. 634 17. 934 NYZRM2001803 42. 649 14. 914 59. 041 11. 657 28. 809 14. 670 17. 172 5. 396 NYZRM2001803 31. 621 19. 917 31. 650 19. 861 14. 683 19. 396 24. 619 18. 912 NYZRM2001804 19. 995 18. 787 31. 384 9. 308 6. 192 7. 945 11. 032 6. 537 NYZRM2001898 5201. 332 1475. 462 2605. 875 738. 729 3013. 651 5911. 225 5347. 627 1306. 593 NYZRM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NYZRM2001902 9. 512 5. 176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NYZRM2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NYZRM2001935 36. 519 23. 148 18. 415 4. 134 11. 165 15. 562 19. 141 10. 042 NYZRM2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NYZRM2001938 23. 961 5. 651 21. 192 6. 301 5. 377 17. 197 7. 025 5. 482										
NTZRM2001865   33. 026   24. 176   29. 953   16. 912   19. 394   23. 562   31. 355   25. 910     NTZRM2001867   30. 838   22. 957   35. 457   14. 948   16. 183   32. 799   17. 562   46. 800     NTZRM2001869   129. 599   162. 083   180. 222   173. 694   64. 737   231. 277   145. 176   147. 129     NTZRM2001879   14. 477   14. 016   20. 104   6. 241   7. 997   18. 463   6. 634   17. 934     NTZRM2001803   42. 649   14. 914   59. 041   11. 657   28. 809   14. 670   17. 172   5. 396     NTZRM2001806   31. 621   19. 917   31. 650   19. 861   14. 663   19. 396   24. 619   18. 912     NTZRM2001806   31. 621   19. 917   31. 650   19. 861   14. 663   19. 396   24. 619   18. 912     NTZRM2001806   201. 332   1475. 462   2605. 875   738. 729   3013. 651   5911. 225   5347. 627   1306. 593     NTZRM2001902   9. 512   5. 1676   9. 030   3. 230   3. 539   7. 418   7. 583   3. 383     NTZRM2001903   63. 243   40. 127   55. 162   28. 793   22. 732   77. 356   28. 595   48. 438     NTZRM2001935   36. 519   23. 148   18. 415   4. 134   11. 165   15. 562   19. 141   10. 042     NTZRM2001936   78. 536   47. 939   51. 879   18. 980   21. 013   48. 576   35. 554   52. 419     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482     NYZRM2001938   23. 961   5. 651   21. 192   6. 301   5. 377   17. 197   7. 025   5. 482		MT2RM2001840								
NT2RM2001867   30, 838   22,957   35,457   14,948   16,183   32,799   17,562   46,800     NT2RM2001869   129,599   162,083   180,222   173,694   64,737   231,277   145,176   147,129     NT2RM2001879   14,477   14,016   20,104   6,241   7,997   18,463   6,634   17,934     NT2RM2001803   42,649   14,914   59,041   11,657   28,809   14,670   17,172   5,396     NT2RM2001886   31,621   19,917   31,650   19,861   14,683   19,396   24,619   18,912     NT2RM2001807   19,995   18,787   31,384   9,308   6,192   7,945   11,032   6,537     NT2RM2001807   5201,332   1475,462   2605,875   738,729   3013,651   5911,225   5347,627   1306,593     NT2RM2001902   9,512   5,176   9,030   3,230   3,539   7,418   7,583   3,383     NT2RM2001903   63,243   40,127   55,162   28,793   22,732   77,356   28,595   48,438     NT2RM2001903   108,255   64,649   109,195   31,339   39,123   80,005   62,289   59,426     NT2RM2001935   36,519   23,148   18,415   4,134   11,165   15,562   19,141   10,042     NT2RM2001938   78,536   47,939   51,879   18,980   21,013   48,576   35,554   52,419     NY2RM2001938   23,961   5,651   21,192   6,301   5,377   17,197   7,025   5,482										
NTZRM2001869 129.599 162.083 180.222 173.694 64.737 231.277 145.176 147.129 NTZRM2001879 14.477 14.016 20.104 6.241 7.997 18.463 6.634 17.934 NTZRM2001803 42.649 14.914 59.041 11.657 28.809 14.670 17.172 5.396 NTZRM2001886 31.621 19.917 31.650 19.861 14.683 19.396 24.619 18.912 NTZRM2001887 19.995 18.787 31.384 9.308 6.192 7.945 11.032 6.537 NTZRM2001887 5201.332 1475.462 2605.875 738.729 3013.651 5911.225 5347.627 1306.593 NTZRM2001902 9.512 5.176 9.030 3.230 3.539 7.418 7.583 3.383 NTZRM2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438 NTZRM2001903 0108.255 64.649 109.195 31.339 39.123 80.005 62.289 59.426 NTZRM2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 NTZRM2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NY2RM2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
NT 2 RM 2 CO 18 B										
NTZRM2001808   14.914   59.041   11.657   28.809   14.670   17.172   5.396   NTZRM2001806   31.621   19.917   31.650   19.861   14.683   19.396   24.619   18.912   NTZRM2001807   19.995   18.787   31.384   9.308   6.192   7.945   11.032   6.537   NTZRM2001808   5201.332   1475.462   2605.875   738.729   3013.651   5911.225   5347.627   1306.593   NTZRM2001902   9.512   5.176   9.030   3.230   3.539   7.418   7.583   3.383   NTZRM2001903   63.243   40.127   55.162   28.793   22.732   77.356   28.595   48.438   NTZRM2001903   63.243   40.127   55.162   28.793   22.732   77.356   28.595   48.438   NTZRM2001903   0108.255   64.649   109.195   31.339   39.123   80.005   62.289   59.426   NTZRM2001935   36.519   23.148   18.415   4.134   11.165   15.562   19.141   10.042   NTZRM2001936   78.536   47.939   51.879   18.980   21.013   48.576   35.554   52.418   NYZRM2001938   23.961   5.651   21.192   6.301   5.377   17.197   7.025   5.482   4.448   34.448										
NT2RH2001886 31.621 19.917 31.650 19.861 14.683 19.396 24.619 18.912 NT2RH2001887 19.995 18.787 31.384 9.308 6.192 7.945 11.032 6.537 NT2RH2001898 5201.332 1475.462 2605.875 738.729 3013.651 5911.225 5347.627 1306.593 NT2RH2001902 9.512 5.176 9.030 3.230 3.539 7.418 7.583 3.383 NT2RH2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438 NT2RH2001930 108.255 64.649 109.195 31.339 39.123 80.005 62.289 59.426 NT2RH2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 NT2RH2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NT2RH2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										1
NT2RH2001887   19.995   18.787   31.384   9.308   6.192   7.945   11.032   6.537     NT2RH2001886   5201.332   1475.462   2605.875   738.729   3013.651   5911.225   5347.627   1306.593     NT2RH2001902   9.512   5.176   9.030   3.230   3.539   7.418   7.583   3.383     NT2RH2001903   63.243   40.127   55.162   28.793   22.732   77.356   28.595   48.438     NT2RH2001930   108.255   64.649   109.195   31.339   39.123   80.005   62.289   59.426     NT2RH2001935   36.519   23.148   18.415   4.134   11.165   15.562   19.141   10.042     NT2RH2001936   78.536   47.939   51.879   18.980   21.013   48.576   35.554   52.419     NY2RH2001938   23.961   5.651   21.192   6.301   5.377   17.197   7.025   5.482     NY2RH2001938   23.961   5.651   21.192   6.301   5.377   17.197   7.025   5.482     NY2RH2001938   23.961   5.651   21.192   6.301   5.377   17.197   7.025   5.482     NY2RH2001938   23.961   5.651   21.192   6.301   5.377   17.197   7.025   5.482     NY2RH2001938   7.88888   7.88888   7.88888   7.88888   7.88888   7.88888   7.88888   7.88888   7.88888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.888888   7.8888888   7.8888888   7.88888888   7.88888888   7.88888888   7.8888888   7.88888888   7.88888888   7.8888888888										
NT2RH2001898 5201. 332 1475. 462 2605. 875 738. 729 3013. 651 6911. 225 5347. 627 1306. 593 NT2RH2001902 9. 512 5176 9. 030 3. 230 3. 539 7. 418 7. 583 3. 383 NT2RH2001903 63. 243 40. 127 55. 162 28. 793 22. 732 77. 356 28. 595 48. 438 NT2RH2001930 108. 255 64. 649 109. 195 31. 339 39. 123 80. 005 62. 289 59. 426 NT2RH2001935 36. 519 23. 148 18. 415 4. 134 11. 165 15. 562 19. 141 10. 042 NT2RH2001936 78. 536 47. 939 51. 879 18. 980 21. 013 48. 576 35. 554 52. 419 NY2RH2001938 23. 961 5. 651 21. 192 6. 301 5. 377 17. 197 7. 025 5. 482										
NT2RH2001902 9.512 5.176 9.030 3.230 3.539 7.418 7.583 3.383 NT2RH2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438 NY2RH2001930 108.255 64.649 109.195 31.339 39.123 80.005 62.289 59.426 NY2RH2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 NT2RH2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NY2RH2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482					DEOK 975					
NY2RM2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438  HY2RM2001903 63.243 40.127 55.162 28.793 22.732 77.356 28.595 48.438  HY2RM2001930 108.255 64.649 109.195 31.339 39.123 80.005 62.289 59.426  HY2RM2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042  NY2RM2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419  HY2RM2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
HY2RH2001930 108.255 64.649 109.195 31.339 39.123 80.005 62.289 59.426 HY2RH2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 HY2RH2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 HY2RH2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
NY2RH2001935 36.519 23.148 18.415 4.134 11.165 15.562 19.141 10.042 NY2RH2001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NY2RH2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
NTZRUZ001936 78.536 47.939 51.879 18.980 21.013 48.576 35.554 52.419 NYZRUZ001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
NY2RH2001938 23.961 5.651 21.192 6.301 5.377 17.197 7.025 5.482										
MIZRUZUU1939 23. 501 3. 031 21. 152 0. 000 100 100 100 100 100 100 100 100										
MIGHTANIAGI   11.430   43.030   10.324   13.140   EE.E.										
		[N1 CMJCD0134]	1 11.430	43.030	1 (0. 323	1 .5. 100	1			<u>.                                    </u>

Table 69

	HT2RM2001950	46, 415	29.816	36.996	18. 559	8.239	39. 347	13. 956	23. 224
	NT2RH2001952	2.871	2.886	10.623	6. 195	0,000	2. 538	1.846	8. 237
5	NY2RH2001976	42.702	29.344	52.698	20.599	18. 125	57.645	24. 197	33.972
•	NY2RM2001982	20. 947	25. 776	25, 162	18. 275	10.576	18.050	9. 191	14.830
	NT2RH2001983	23.643	16.045	27.661	9.316	13.749	23.964	18. 258	17.035 44.130
	NT2RM2001984	147.043	51. 662	81.658	22.066	35.725 24.785	120. 259 39. 074	90. 102 34. 205	61. 176
	HT2RM2001989	76. 106	50.939	80.150	44. 331 23. 083	19.109	45. 858	26. 865	31. 923
	NY 2RH2001996 NT 2RH2001997	37. 798 63. 158	41.931	43. 246 28. 543	20.691	22.046	58. 320	33.747	34.764
10	NT2RM2001998	47. 869	29.374	50. 969	17.042	23.450	45.674	22.546	20.062
	NT2RD2001999	23.045	23. 925	34, 107	16. 137	19.923	26.601	19.613	27.167
	NT 2RH2002003	60. 554	45. 534	133.518	30.271	23.148	57.270	34.688	36. 304
	NT2RH2002004	16.782	14.896	24. 193	8.483	9.918	13.788	12. 592	4.939
	NT2RH2002009	22.784	26.292	37.573	16.205	17.990	24. 047	10.371	17. 159
15	NT2RH2002014	12.027	11.499	20.605	9.676	8.686	10. 127	8.085	17.091
	NT2RM2002019	45.009	49.617	61.370	29.641	24.044	45. 990 90. 878	20. 852 44. 848	29.924
	NT2RH2002029	100.329	58. 955	73.738	25. 096 23. 542	36.513 18.217	49.856	26. 265	32.557
	NT2RM2002030	53.030 55.319	36, 122 58, 655;	48.637 69.310	15.775	34. 969	119.355	37. 851	31.637
	NT2RM2002034 NT2RM2002049	30.306	26. 333	67. 224	12.461	13.486	32.196	19, 763	26. 143
	NT2RH2002055	4.746	9. 322	10.601	1.587	3.475	2.738	4, 711	1.253
20	HY2RM2002072	274.106	142.825	221.668	99.170	111.051	240. 179	193. 919	147.089
	NT2RM2002088	66.101	43. 548	67.009	20. 108	29.769	- 38. 434	34, 203	37.710
	NT2RM2002091	157.752	95. 255	103.301	42.530	51.107	91.971	64.745	58. 827
	NT2RH2002100	36.481	42.661	83.563	34. 382	22.604 17.601	42.960 56.066	30. 266 33. 542	45. 203 37. 167
•	NT2RH2002109	65.961	25.178	54.629 244.199	11.426 79.521	110.685	272. 182	195. 547	168.748
25	NT2RM2002126 NT2RM2002128	271.768 30.978	20. 989	35.773	13.699	15. 221	20. 446	37.022	29. 430 -
	NT2RH2002129	53.911	38. 709	50.544	14. 507	24.022	54. 089	40. 427	15.416
	NT2RM2002142	157.794	95, 271	127.900	44. 871	54.994	121.896	116.748	122.762
	HT2RM2002144	39.141	23.769	42.061	18. 362	18.425	90. 424	32.619	22.086
	NT2RM2002145	69.465	33.538	54.629	19.065	28.804	64.861	31.013	26. 312
20	NT2R12002153	57.982	34.658	45.808	37. 204	22.363	85.615	33. 858 18. 529	45. 468 24. 578
30	HT2R12002163	46.164	22.611	32.853	10.533	12.313 7.659	28.767	6.584	31.812
	NY2R12002170 NY2R12002178	20.367 72.826	15. 918 29. 934	26. 954 35. 113	17.854	17.814	57.676	53.788	36.064
	NT2RH2002179	20.487	16.890	26.778	4. 596	7.536	27. 483	11.691	22.514
	HT2R/2002270	75.965	30. 835	59.481	19. 162	23.264	67.579	38.824	31. 179
	NT2RE2002326	25.054	17.109	25.901	10.631	13.295	20.170	15. 155	11.219
35	HT2RM2CO2337	49.608	30, 430	44. 382	14.424	20.214	49.783	38.536	36.266
	HT2RM2002339	126.783	46.855	62.446	22.680	35.280	129.046 27.503	67.853 20.940	46.026 24.302
	NT2RH2002345	34.662	27. 251	30.489 118.627	17.636 55.152	9.930 36.416	61.876	35. 957	79. 909
	NT2RE12002368	53.018 29.049	67. 271 17, 380	20.968	5. 965	9.584	35.715	13.371	27.731
	NT2802002424	23.738	30.901	58.344	39. 153	17.434	49.766	25. 216	77.325
40	NT2RM2002450	40.370	29. 535	54.082	14. 242	16.219	34.988	19.676	33.464
	NT2RM2002482	44.705	26.737	46.955	14.769	18.437	42,664	46.045	30.188
	HT2RM2002492	113, 197	127. 579	109.738	72.932	49. 321	103. 335	74.905	97.173 62.455
	MY 2RM2002575	112.457	88.605	247.074	59.323 57.513	48.212 25.109	80.685	30.240	69.813
	NT2RM2002580	64.838	62.853	96, 103	45. 340	44.856	104. 438	69.434	96. 173
	NT2RM2002592	20, 462	70. 152 48. 581	29.949	14. 231	13.430	29.384	17.823	61.212
45	NT2RH2002615	33, 564	24. 375	25. 868	12.468	16.085	46. 176	71.059	33.280
	NY2RM2002522	95.365	53.669	62.071	44. 205	38.612	108.504	47.073	91.258
	NY28M2002630	118.784	86.444	276.792	68.615	58.079	85.846	51.946	80.285
	NT2RM2002634	36.887	30.749	31.925	22. 948	20.353	42.111	32. 736	22.117
	HT2R12002645	51.215	209.069	58. 292	23. 942	32.501	97.660 50.329	24.132	61.537 24.267
50	NT2RH2002646	69, 318	57.452	61.629	25.645				42.269
	NY2R12002647	31.140 42.576	27. 535 30. 866	50. 514 34. 782	14.850			14.955	30. 578
	HT2RH2002652	53.871	40.724	63. 208					
	NY28112002721	81.740	78. 721	123. 105			98, 931	44. 593	
	NY28112002748	91. 982	206.064		241.969	54.156	135.810		
55	HT2RH2002764	46.071	41.769				34. 165	32.761	36.777

## Table 70

	NT2RH2002772	80.296	40, 944	68. 101	23.056	28. 389	72.818	41.505	60. 302
	NT2RM2002811	63.439	38.909	43.044	17.983	20. 375	56.523	23.815	28. 434
	NT2RH2002818	50.605	52.430	151.915	32.193	19.702	26.680	17. 380	40.512
5	NY2RH2002879	24, 562	28. 586	34. 172	8.860	5.095	18.514	12.159	30.354
	NT2RM2002979	84. 387	41.192	53.776	21.436	31.083	74.067	53.736	47.429
	NT2RM2002981	59. 340	25.706	33. 191	11.478	15. 597	54.899	35. 830	32.861
	NT2RH2002995	42.179	21, 303	31.267	13.206	10.830	32.109	30.448	42.538
	NT2RU2003031	44.114	29, 430	46.063	16.774	17. 437	43.222	40.155	25.053
	NT2RM2003042	106.509	160.917	155. 488	83.058	73, 174	152.473	69.308	122.583
10	NT2RM2003044	33.909	33, 603	47.142	12.698	45. 517	25.310	25. 508	29. 529
	NT2RM2003090	47.953	25. 520	41.051	9,604	15. 180	34. 197	23. 552	25.659
	NT2RM2003095	43.943	31.580	32.103	11.759	18. 398	29. 592	34. 566	28.874
	MY2RM2003116	20.590	18.126	22.701	10.734	10.194	11.727	12. 203	14.479
	NT2RM2003222	21.398	10.313	27.148	5. 349	13.395	13.068	20. 550	25. 145
45	NT2RM2003224	110.266	37.406	48.819	30.835	29. 947	80. 454	57.677	53. 588
15	NT2RM2003250	30, 062	26.498	38.776	15.773	16.547	23.997	24.660	26.915
	NT2RM2003258	12.707	12.077	15.752	5. 247	7.979	8. 239	5.752	8.852
	NT2RM2003262	37. 575	42.567	50.603	27.374	33.378	31.965	36. 375	43.803 35.943
	NT2RH4000023	49.690	44.882	57. 421	17. 352	24.868	53.007 34.975	25. 083 25. 466	17.156
	NT2RH4000024	33.710	23.142	26.564	7.803	10.308 3.969	5, 783	1.681	9. 230
20	NT2RM4000027	6.576 107.340	5. 402 43. 649	9. 541 64. 579	2. 488 25. 595	27. 984	81.398	45. 801	45.851
20	NT2RD4000033	54. 521	41. 188	116.087	19.883	18. 324	28. 028	14.764	29. 244
	NT2RH4000034	8.646	20. 135	21.495	9.212	9.086	13.100	7. 920	12.176
	NT2RD4000046	42.055	17.446	23.148	8. 687	9. 540	32. 532	23.736	18.823
	NT2R14000052	23.740	17. 236	25.146	8.065	5. 341	17.707	13.080	13.561
	NT2RM4000054	440. 502	221.475	352.643	107.153	132. 322	410.274	281.112	209.475
25	NT2RM4000051	30.264	15.792	27.807	6.396	10.845	21.557	14. 902	4. 276
	NT2RM4000074	8.073	35. 126	41.073	20.510	9. 480	34. 431	24. 493	47.368
	HT2RM4000085	22.897	19.315	23.277	16.541	12.977	24. 111	12. 451	24.618
	NT2RM4000086	50.715	22.670	78.725	20. 299	18.217	28.085	16.663	27.361
	NT2R14000100	17.872	21.935	15.019	10.707	10.091	15. 556	12. 260	12.129 5.064
	HY2R14000101	42.770	15. 330	25.674	6.552	7.785 208.613	24. 576 334. 316	212.009	231.229
30	NY2814000102	407.848	190. 329	321.537	152.733 3.131	7, 950	21. 156	10.845	7.969
	NT2RM4000104	23.885 32.088	13.626 10.072	17.310	5. 693	9. 226	13, 512	10.582	7. 588
	NT2RM4000113	36.681	21.490	22.965	12.521	11.849	23.308	16. 146	10.761
	NT2RM4000139	25. 930	23.620	31.564	24.607	22.610	18. 556	14.008	44.620
	NT2RM4000149	33.404	17. 925	29.734	13.712	15. 989	18.474	26.736	42.075
35	NT2RM4000155	21.566	44. 820	46.750	15.598	16.524	14. 928	9.733	8.224
33	NT2RE4000156	16.586	6. 239	5.822	3.387	3. 958	28. 594	7.207	15.119
	NT2RM4000167	20.171	16.879	15.859	11.667	2, 739	8. 443	3.474	21.050
	NT2RH4000169	30.428	28.089	36.443	24. 244	11.338	20. 566	13. 227	60. 152
	NT2RM4000191	52.656	25. 321	40.946	12.980	18.787	41.092	35.047	38, 394
	NT2R24000197	15. 240	11.946	16.612	2. 282	13. 434	15. 387 49. 371	8. 823 53. 195	5.757 32.774
40	NT2RM4000198	88. 525	63.904	196.728 46.280	39.099 17.110	18. 960	33. 287	27. 322	30. 945
	NT2RM4000199 NT2RM4000200	52.380 33.395	16. 462	28.537	10.600	16, 103	20.714	14.030	6. 949
	NT2RU4000202	30, 208	20. 922	42.468	9. 182	9. 970	16. 908	10.274	12.811
	NT2RE4000210	66.407	27.815	30.474	15. 335	16.812	41.212	27.389	47. 172
	NT2RM4000215	25.869	24.845	36.251	22.848	13, 152	31.488	12.403	27. 548
	NT2RM4000220	47. 201	39. 573	38.877	20. 267	19. 583	51. 592	35. 424	51.912
45	NT2RE4000229	38.395	26.396	42.302	13.878	14. 171	29.316	28. 242	16.590
	NT2RE4000231	54.697	33.959	43.440	18.016	23.895	29. 537	28. 746	34.406
	NT2RM4000233	209.479	90. 187	137.270	36. 159	66.994	160.853	100.732	62.965
	HY2RH4000244	16.916	9.010	13.401	4. 357	9.911	12. 907	8.771	8.963
	NT2RH4000251	43.833	19.474	33, 500	11.060	16.673	31.966	32.833	8. 105
50	NT2RM4000255	35.799	17. 398	36.446	10.625	12.098	. 29. 741	23. 847	15. 929
50	HT2RE4000265	102.046	79.778	222.138	64.769	51.026	72.136	39.083	49.420 166.824
	MY2804000283	285. 571	172. 391	189.067	109. 857	94. 953	255. 306 25. 501	162. 352 27. 248	34. 927
	MT2RM4000284	23.615	36. 279	30. 562	12. 441	17.835 22.008	73. 912	45.709	43. 178
	NT2RM4000290	74.673	36.513 18.871	57.081	15. 623 8. 987	11,022	47. 890	18.701	14.976
	NY2RM4000295	24.000		22.693 51.817	18.306	78. 561	140.750	92.030	34. 220
55	NT2RM4000306	140.029	42.148	1 91.011	1 10. 300	10.301	1 170.700	1 32.000	*****

Table 71

	NY2RH4000307	20.678	19. 168	22.141	9.050	9. 145	23.385	14. 343	13.754
	NY2RH4000309	41.662	20.618	26.408	8. 581	10.787	30.894	18.116	11.868
	NT2RH4000313	35.434	20. 403	33. 260	17.080	12.239	39. 520	34.145	43.040
5									
•	NT2RM4000318	52. 262	31.467	139. 471	20.774	17.880	23.820	17.441	19.608
	NT2RM4000324	51.333	27.748	39.958	9.932	17. 995	63.248	27.625	42.800
	NT2RM4000326	32, 179	16.471	20. 536	8.435	10.621	23.791	17.926	20.620
	NT2RM4000327	60. 230	58.958	198.666	39. 302	28. 376	44.008	20.961	43.734
	NT2RH4000344	63.708	65. 489	173.360	38. 949	27. 536	34. 270	15.519	42.106
	NT2RM4000349	30.022	14.663	14.070	7.442	10. 197	22. 535	12.455	16.210
10	NT2RH4000354	46.698	15. 085	27.013	11.329	7.922	27.895	13.694	15.005
	NT2RH4000356	32.497	24. 336	32.372	13.972	11.464	43.673	31.608	29. 630
	NT2RH4000366	528. 262	330.865	423, 109	167.985	170.232	378.411	215.606	442.307
	NT2RH4000368	51.220	51.300	153. 236	33.445	22.538	43.253	17. 539	64.383
	NT2RM4000373	25. 297	22, 861	32.020	19.516	16.128	25.045	13.784	37.614
	NT2RM4000386	22.576	9.738	24.078	8.987	9.704	21.730	24.414	23.758
15	NT2R14000395	61.364	79.696	124.563	37.133	40.433	107.248	46.227	46.047
	NT 2RM4000414	159.474	59. 130	69.911	18.566	40.333	119.002	79.051	21.561
	NT2RM4000417	15.712	20.634	23. 502	7.213	7. 502	15.030	7.412	1.867
	NT2RM4000421	15. 106	14.708,	19.062	8.549	6.469	15. 114	8.074	20. 588
	NT2RM4000425	101.441	83.854	259. 486	55, 511	39.319	53.250	31.739	69.026
	NT2RM4000433	51.457	24.650	39.654	12.379	16.608	41.763	37.139	36.708
20									
20	NT2RM4000436	51.207	21.755	29. 107	13.444	12.333	34.290	27.223	37.320
	NY2RM4000444	40.864	26.268	67.826	11.797	17.600	39.060	23.113	28.672
	NT2RM4000457	63.983	39.080	61.124	23.292	28.748	50.040	26.813	31.965
	NY2RM4000471	41.652	29.088	37.803	8, 939	15,093	35. 469	20.877	14.796
	NY2RM4000472	68.502	62.226	206.357	48.752	23,646	77.597	28. 412	104.099
	NT2RM4000486	30.140	26.427	28. 452	18.097	7. 542	22.184	12.697	24.533
25	NT2RM4000490	51.124	23. 641	42.235	9, 300	14.683	56.785	25.625	17, 105
	NT2RH4000496	110.770	31.642	65.060	13.739	27.500	68.720	52.247	37.631
	MT2RM4000505	134. 100	84.063	126.035	43.665	56.053	130.720	81.120	71.520
	NT2RM4000511	73.441	160.671	81.146	172.018	35.906	98. 128	55.037	164. 299
	NT2RM4000514	24.804	23.670	34.085	13.945	16.589	32.103	21.758	11.170
	NT2RE4000515	56.528	99.798	88.516	40.030	41.279	67.061	40.210	72, 202
30	NT2RE4000517	94. 295	97.384	143. 107	76.451	43. 905	144.940	69.520	145.604
	NT2RM4000520	13.459	13.780	16.902	5. 273	5. 564	7.899	7.054	14.968
	NT2RM4000531	29. 188	24. 283	26.738	11.063	12.826	18. 929	23.443	20.712
	NY2RM4000532	14. 395	12.711	19.277	9. 437	8. 520	12.914	15.215	13.835
	NT2RM4000533	18.380	13.704	18. 165	8. 534	7.454	15.515	10.288	7.686
	NT2RM4000534	17.803	11.768	18.975	7. 585	10.236	14, 119	11.420	19.497
35	NT2RM4000563	53.983	34.056	51.401	17.700	36.352	45.609	32.373	33.367
33	NT 2RM4000566	36.586	22.989	35.859	9. 957	21.078	25.668	24.949	21.224
	NT2RH4000568	59. 423	29. 845	36.652	12.139	25.850	70.617	54.001	29. 192
	NT2RM4000585	48.810	27.673	38.443	12.701	20.510	33.948	23.868	27.346
	NY2HMA000587	29.705	26.644	25, 876	12.729	11.927	16, 240	17.926	19.718
	NT2RH4000590	32.164	21.289	29.186	8. 941	11.617	18.856	16.495	13.544
	NY2RM4000593	61.080					33.032	30.484	25.715
40			32.766	38.970	15.411	20.360			
	NT2RE4000595	41.141	22. 473	35.313	9.766	11.448	11.237	20.012	12.069
	NT2RM4000603	78.976	52.410	58.176	24.839	24.042	50.072	40.363	31.910
	NT2RM4000611	15.953	10.734	13.469	9.013	8.977	10.161	7, 157	22.979
	MYZRE4000516	45.814	37.309	35. 175	17. 505	23.768	40, 117	27.918	39,007
		43.014							
	NT2RM4000621	57.493	77.709	73.014	76.819	24.081	71.204	46.769	83. 169
45	NT2RM4000648	28.637	18.518	26.908	8. 210	13.083	15.965	12.644	11.022
45	MY2RM4000649	85.058	41.743	59,668	13.629	29.612	55.983	39.586	36.405
	NT2RM4000658	135.688	61.028	120.722	28. 197	43.765	79.777	46.011	96.630
	NT2RM4000561	71.854	99. 345	52. 294	18.409	29. 132	62.897	45.030	41.904
	NT2R04000673	135.680	61.584	75.017	24. 321	20.618	70.048	46.608	45. 107
	NT2RM4000674	75.722	36.633	51.480	16.765	16.961	34.561	42.749	30.664
	NT2RM4000889						22.615	15.641	20.045
50		41.790	28.540	39.966	15. 401	8. 448			
	NT2RD4000598	61.169	46.347	64.951	24. 102	41.257	63.885	38.390	29.637
	NT2RM4000700	27.239	106.106	27.114	9. 273	11.699	12.813	14.815	12.082
•	NT2RM4000701	227.264	115.040	182.483	47, 970	70.324	76.813	128, 958	65, 330
	NT2RD4000712	43, 183	27.951	46. 394	10.240	14, 368	19.562	26.208	16.644
	NT2RM4000717	34. 386	22.333	19. 262	10.038	12.975	19.299	13.148	20.540
	NT2RD4000733	75.958	43.996	58. 928	24.743	28.885	88.871	65.331	37, 193
55	···	1 13.330	1 -5. 550	1 55. 550	1				

Table 72

	NY2RH4000734	24. 197	38. 270	53. 725	16.970	13, 155	39.087	23.333	39. 227
	NT2RH4000741	43.844	13. 589	30. 427	10.346	8.744	26.119	12.592	26.083
	NT2RM4000744	50. 833	14. 548	25.024	23.480	10.805	62, 136	17.742	83.553
5									
	NT2RH4000749	80.902	71.083	91.633	27.354	60.031	198.030	52.328	100.669
	NT2RM4000751	22.688	29.768	53, 788	53. 315	27. 282	19.811	22.272	42.714
	NT2RH4000752	52. 247	32.866	40.812	14. 427	15. 224	9. 355	23.407	43. 927
	NT2RM4000760	33.235	16. 169	27. 997	11.989	19.412	13.254	10.563	10.820
	NT2RM4000761	2403.264	848. 134	3887, 956	172. 265			2359.029	400.128
	NT2RE4000764	301, 709	144. 132	163.494	49.659	143.743	257.369	245.639	103.045
10							10.345		11, 267
	NT2RH4000768	11.747	9. 247	11.542	9.135	9. 038		6.336	
	NT2RH4000778	6.893	5.725	9. 950	5. 466	4. 458	5, 886	5.079	5.685
			96. 516		51.850	99.170	184.671	138. 565	75.926
	NT2RH4000779	238. 073		182.851					
	NT2RH4000787	69.121	57. 977	157, 708	28. 426	29. 213	21.609	22. 533	11.420
		60. 309	45,026	83. 182	23.988	30, 494	22.815	35. 485	31.417
	NT2RH4000790								
4.5	NT2RM4000795	453. 425	108, 548	204,710	17.809	92. 365	272.802	147.653	47.088
15	NT2RE4000796	144. 288	57.098	70.720	23.213	47, 104	97.550	50, 426	30.942
									11.548
	NT2RM4000798	59. 938	28. 301	25.839	10.244	18. 327	23.444	20.572	
	NT2RH4000800	150.768	122.487	195.880	137.376	57. 284	146.130	97.369	185.386
					12.655	14. 527	25. 975	22.848	11,921
	NY2RH4000813	37.084	20.876	36.294					
	NT2RU4000820	86.855	60.381	192.196	39.751	37.738	50. 427	35.797	26.747
	NT2RM4000827	41.788	28.006	51.622	20.945	21.631	21.541	30. 438	31.570
20								30. 029	44, 496
	NT2RM4000830	68. 078	30.965	59.647	20. 203	26. 347	37.484		
	NT2RH4000833	111, 407	74.480	77.732	17.832	39. 802	56.697	25. 292	36.404
	NT2RL4000841	49, 942	45. 599	72, 313	16.308	20.094	29, 644	26. 188	28.854
									14, 948
	NY2RH4000846	104. 561	76.278	275. 932	57.490	49. 037	63.058	35.772	
	NT2RH4000848	125. 196	36.830	101,007	17, 584	32.806	82.740	51.262	19. 922
	NT2RM4000852	113.009	77.800	126.639	43.464	43.880	57.479	52.365	44. 156
25 ·									51.332
25	NT2RH4000855	64.608	50.229	146.326	22.844	23.661	28. 928	25.813	
	HT2RM4000859	24, 418	19.759	24, 141	10.385	14.916	34. 345	18.598	11.625
	NT2RM4000868		14.752	14, 556	11.565	9.114	12.226	17.324	12.029
		15. 564							
	NY2RM4000870	55, 531	47.020	57.796	18.791	30. 154	39.778	25. 127	26.057
	NT2RM4000879	103.887	41.773	56.495	12.837	31, 154	67.942	43.586	22.044
	NT2R04000882	81. 982	42.561	80.304	22.840	38,713	36.853	45.646	48.992
30	NT 2RM 4000887	151.731	36.758	112.092	22.645	40.960	98. 527	85. 229	22.008
	NT 2R14000895	84, 679	41.293	172. 935	28.755	27.724	44. 297	19.644	26. 291
				58. 329	17.578	25. 299	44.317	41.019	30.575
	NT2RM4000897	45. 994	42.630						
	NT2RU4000901	13, 138	13.528	18.046	7.930	5. 669	7.738	9.304	5.798
	NT2RM4000950	13,710	21.028	17, 402	10.585	11.390	13.090	8. 272	13.397
							23.064	21.414	26.049
	NT2RM4000965	54. 459	36.282	50.127	15.952	25. 327			
35	NT2RB4000971	41, 258	27.847	39.604	12.433	17.061	72.230	20.025	17.430
	NY2RH4000979	33. 580	21.677	32.692	7.475	11.647	22. 259	16.549	12.389
							42.714	19.808	18.064
	NT2RM4000987	51. 537	23.981	27.883	11.309	12.974			
	NT2RM4000989	43. 246	16.680	33.780	10.504	10.430	22. 581	33.282	15. 259
	NY2RH4000991	6. 595	8.954	14.910	4.216	4,093	24. 193	3.472	15. 581
							33.667	22.750	38, 582
	NT2RM4000992	61, 901	44.659	179.747	37. 376	29. 327			
40	NY2RH4000996	12.902	17.829	47.104	22.304	9. 589	15. 133	12.379	41.017
	NT2RM4000997	139.754	107.958	216.478	45.750	59.135	79.871	47.855	52.159
				123.641	25. 902	74.114	102.439	120.879	88.667
	HT2RH4001001	222, 229	90.117						
	NT2RH4001002	22.453	23.223	34. 127	15.841	13. 942	17.616	10.393	26.669
	NT2RM4001016	39. 433	22.372	27.844	7.677	15, 230	29.791	22.346	14.840
				262.665		89.809	167.042	104.628	258. 452
	NT2RM4001025	123, 159							
45	NT2RM4001027	1,003	0.083	0.000	0.188	1. 139	0.903	0.000	13.341
45	NT2RM4001032	15.446	8.560	20. 283	7.827	10.702	9, 129	9.798	10.321
						7. 503	4. 130	9. 323	18.916
	DT2RU4001047	18. 565	7. 922	16.869	2.924				
	NT2RM4001049	87. 157	64.640	99.050	20.618	35. 192	44. 265	24. 923	27.816
	NT2RM4001051	45. 597	65.440	63.291	17.761	11.312	31, 198	20.661	24. 356
					12.670		36.706	54.060	39.934
	MT2RM4001052	83.704	54.084	58.884		16. 509			
	NT2RE4001053	55. 548	69.868	192. 178	27.160	24.862	42.613	24. 525	28.003
50	HT2RM4001054	29, 223	12.533	27.929	5,313	10.023	15, 125	15,911	14. 263
,									64.703
•	1012RE4001059	181.587	40.368	91.633	17.857	33.606	105. 399	88.210	
	NT 2 REMODITO Y	29.020	21.136	81, 470	8.928	13.093	5.999	16.142	11.555
				39.816	12.808	14. 924	24, 390	23.123	13.779
	MT2RM4001084	42.690	28. 922						
	NT2RM4001092	102.531	57.027	86.268	31.684	25.916	49. 946	45.616	58. 081
	NY2854001100	43. 266	33.448	49.943	8.293	19.072	24.126	16.221	46.701
55	FULL CUT-MODILION	1 43.200	30.770	77.373	1 5.23	1 .3.0.2			

Table 73

HT2RH4001116	27.726	26.051	28. 521	6.793	9.001	18.038	14, 406	8. 177
NT2RH4001119	56.668	21.890	35. 980	9.796	15, 859	38. 916	35. 588	15.608
NT2RH4001140	136. 817	79.720	322.522	72.609	64. 281	53.073	51.451	56.047
NT2RH4001148	238. 824	52.972	84.009	16. 224	62. 535	137.805	147.073	38.797
NT2RM4001151	49.119	18.810	31.963	9, 013	16.522	24. 362	37.118	17.496
NT2RM4001155	51.322	26.524	38. 663	9.832	19, 192	16.401	24. 191	12.958
NT2RM4001157	29. 926	19. 538	29. 560	8. 442	11.794	23.764	9. 393	5.071
NT2RM4001160	72. 399	50. 574	60.230	13. 285	29. 392	49.862	35. 181	33.807
NT2RM4001163	150.688	70, 942	95.070	47. 204	58.092	77.447	65. 645	40.117
NT2RM4001187	46.613	33. 566	37.323	10.669	19.756	22. 493	19.909	13.410
NT2RM4001191	62. 821	78. 568	138. 398	23.085	37. 250	19.851	28.068	31.505
NT2RM4001200	48. 487	41.856	115. 958	43.120	35.674	29. 433	29.755	46.933
NT2RM4001203	29.740	33. 257	26. 183	10.711	18.414	17.515	13.820	29.510
NT2RM4001204	85. 368	2.729	5. 406	1.939	1.539	2.503	5. 782	1.987
NT2RH4001217	22.326	14. 483	20.894	6.910	10. 252	17.142	14, 178	16.377
NT2RM4001245	102.964	61.341	59. 224	17.873	32. 130	47.902	39.713	28.855
NT2RM4001247	60. 472	48. 248	105.685	27.869	20.131	20.633	22.912	17.998
NT2RM4001256	38. 132	20.867	27. 791	11.662	11.297	22.362	18. 443	14. 221
NT2RM4001258	13. 173	14. 508	15, 622	2.115	6.064	10.903	11, 147	31.184
NT2RM4001267	18. 994	10.887	19.555	6. 271	8. 494	3.421	7.779	13.809
NT2RE4001273								39.740
	57. 388	34. 293	59.413	25. 522	17.714	21.978	30.691	
NT2RM4001281	52. 686	24. 825	33.241	13.708	11.390	31.923	19, 522	23.080
NT2RM4001286		1240. 433	782.259	477.895	296.841	681.688	413.930	936.577
NT2RM4001290	25. 298	23.154	13.373	6. 552	0.000	12.469	8.723	14.611
NT2RM4001309	48. 445	24.031	36.511	15.060	18. 354	33.040	18, 409	21.487
NT2RM4001313	61.618	55. 950	171.030	27.704	18. 541	31.137	15. 527	37.397
NT2R04001316	49. 175	40. 348	93. 903	19.571	16. 907	28. 903	20. 127	14.212
		40. 346						27.654
NY2RM4001320	73.145	43. 895	149.769	28. 755	24. 031	24. 203	22.793	
NT2RM4001321	49. 367	26.564	28. 912	10. 370	15. 275	21. 145	21. 285	20.579
NT2RM4001325	38. 855	43. 433	53. 158	15. 234	25. 333	31. 524	26. 184	15.840
NY2RM4001333	48. 466	17, 343	99.002	20. 144	115, 167	148.955	12, 312	8, 170
NT2RM4001340	30.804	28. 992	40. 576	27.062	32.009	10. 155	18.551	26.573
NT2804001344	30.624				12.525	9. 910	11.004	11.417
		35.092	33, 290	12.667				
NT2RM4001347	14. 549	14.691	20.853	11.657	13. 229	14, 366	8. 959	54.748
NT2RM4001357	58.256	26. 925	40.009	14.812	13.213	104.908	348.697	7.592
NY2RM4001360	86.062	33.099	53.959	12.261	27.140	48. 858	36.604	20.008
NY 2 REM 4001371	57.075	37.841	49.730	24. 239	25.868	54.098	8.910	31.242
NT2RM4001377	101.216	75, 138	68.626	19.407	36. 169	52.589	30, 583	31.839
MT2RM4001382	56.509	78. 201	56. 186	36.607	24.700	70. 227	41.803	66.511
MICHEMOTIOS								
NT2RE4001384	13.506	11, 432	7. 793	6. 199	7. 970	12.881	6.788	7.108
NT2RU4001400	21.837	16.958	21.913	10.795	7.913	16. 255	9. 524	12.188
NT2RM4001409	28. 309	17.011	26.656	9.796	12.960	23.632	14.054	20.949
DT2RM4001410	29.072	19.001	30. 576	8. 925	14. 550	18.489	21.014	17.448
NY2RM4001411	8. 505	7.030	30.358	2.388	3. 324	0.962	1.969	1.931
NT2R14001412	59, 413	25. 935	59.821	15. 231	22.577	30.927	24.563	11, 190
HY2RE4001414	64.093		33.046	9.873	26. 265	24. 538	20.805	20.958
		33, 321						14. 185
NT2RM4001436	33.680	29, 671	20.088	7.331	12.620	14.939	11.458	
NY2RM4001437	70.569	41.529	158.116	28.707	19. 302	25.565	23.649	23.787
NT2RU4001444	63.099	33.815	51.190	21.250	36.920	56. 421	41.830	35.180
NT2RM4001454	15.293	16. 251	33.213	14.589	11.226	13.235	7.237	9. 931
NT2RM4001455	8.536	7.947	12.910	5. 235	6.864	7.007	13.432	28.743
NY2RM4001483	74.168	64. 931	192.825	43.272	33.854	44.722	22. 451	46.563
							23. 221	19. 361
NT2RM4001489	27.884	28, 159	36.108	13.377	14.505	15.628		
HT2RM4001495	260. 493	117. 396	133.602	31.705	64.659	91.833	54. 255	51.382
NT2RM4001499	68.936	37. 210	73.295	19. 265	26.638	41.151	25.000	25.754
NT2R04001515	11.646	7.906	18.332	5.318	7.167	15.640	6.612	8. 512
NY2RM4001519	12.556	9. 937	20.664	5.346	32.689	10.138	7,966	8. 328
HT2RM4001522	71.440	69, 438	164.718	40.425	35, 841	32.755	19.774	38.742
							19.569	31.077
MT2RM4001523	24.710	16.532	29.750	8.848	11.883	12.279		
MY2RM4001550	24. 908	22.060	34. 537	19, 909	20.432	20.143	15. 284	28.090
NT2RM4001553	73.682	40, 371	52.795	27.094	23.686	46.848	27.034	27.166
NY2RM4001554	53.585	30.046	33.134	23.878	15. 283	26.877	16.771	20.649
NT2RE4001557	19.423	19. 434	24. 184	11.971	12.237	21.486	7.653	15. 404
NY2RM4001565	65.552	37. 852	90.440	18.538	17.294	23.128	23.413	18.529
[121 TUT-100   202	1 03.332	1 31.032	30.440	10. 330	11.434	20.760	1 20.713	

Table 74

				1 0 0	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,				
	NY2RM4001566	100.946	48.659	87.457	28.565	28.860	79.976	52.286	9. 785
	NT2RM4001569	7.010	5. 598	41.076	3. 288	8. 597	0. 901	6.611	1, 304
			24.859	37.584	7. 247	15. 119	35, 411	21.050	31.905
5	NT2RM4001579	41.258			10. 109	10.956	22.015	19.971	25. 442
	NT2RM4001582	36.827	23. 162	29.372					35. 619
	NT2RM4001589	57. 574	32.795	61.841	23.877	20. 226	47. 320	41.167	
	NT2RM4001592	32.950	21.429	32.007	7.221	14. 392	17.425	7.965	10.850
	HT2RM4001594	55. 970	26. 805	46.827	13.556	21.275	46.488	34.751	25.706
	NT2RM4001597	113. 189	66.565	189. 284	36.307	35.658	51.457	41. 254	42. 293
40	NT2R54001605	16. 347	11.985	18.084	2.805	4. 141	11.032	9.672	10. 297
10	NT2RE4001609	173.865	587.184	265.155	76.761	120. 584	182.319	73.643	191.832
	NT2RM4001610	89.090	32. 924	55.024	13.942	38. 114	56.107	36.218	28. 535
	NT2R04001611	30.709	14. 204	28.060	6.394	11.242	12.351	22. 333	13.486
	HT2RM4001618	77. 313	59. 231	178, 569	26, 795	28.633	44.101	23.934	50. 341
	NT2RD4001622	42. 484	50.813	37.378	16, 153	35, 073	39. 451	29.062	30. 213
	NY2RM4001624	55. 088	36. 243	39. 342	10.093	11. 389	25. 162	26.300	19.356
15	HT2RM4001625	165. 457	44. 283	55. 076	16.243	29.704	87.349	62.707	32.707
	NT2RD4001629	23. 424	34.729	31.319	10.721	9, 407	17.262	17.006	17, 599
			105.740	108. 162	80. 539	33.853	62.834	39, 339	102. 299
	NT2RM4001632	49. 318		25. 229	7. 187	11.536	12.746	15, 743	16.315
	NT2RM4001642	26.758	24.864	257.397	53, 466	49.798	64,749	33.054	65. 546
	NT2RB4001647	140.643	83.479				10.663	14. 969	25. 969
20	NY2RM4001650	20.039	17.016	26.536	7.633	8.417	43, 545	39, 576	18.233
	NT2RM4001662	93. 433	61.261	62.868	18.713	28.801		21.822	28.747
	NT2RM4001666	99. 250	58. 594	135.514	19.947	25.792	43.075		44. 557
	NY2804001670	108. 596	50.059	60.195	8.757	26.897	80.647	55.639	48. 902
	NY2R24001682	23.010	37.857	52.107	34.229	26.474	24.078	19.040	
	NT2RD4001710	71.974	22.009	43.652	12.553	17. 193	33.805	36. 338	25. 346
	NT2RM4001712	30.145	17.963	29.768	6.775	12.959	13.705	17. 401	11.444
25	NT2RM4001714	39. 284	71.253	45. 168	23.590	23.852	34.014	32. 992	44. 464
	NT2R04001715	39.876	47.568	68. 485	29.814	28.676	29.317	23. 694	38.125
	NT28M4001727	18.826	16.671	24.630	8.765	12.534	14.525	9. 624	7. 446
	NT2854001731	163.786	60.747	103.744	21.266	23.073	109.348	70.159	88.870
	NT2894001735	25. 147	42.977	27.836	33. 257	23. 484	16.531	22.623	48. 984
	NT2804001739	29.621	22.031	33.503	11.627	16.721	10.593	7. 382	14, 863
30	NT2854001741	117.616	80.979	99.834	34.861	34.797	49.703	68.739	91. 553
	NY2R24001746	61.847	44.910	113.561	21.148	31.787	37.464	33.824	23. 274
	NY2894001754	72.161	34.709	70.656	13.473	25. 420	34. 023	22. 194	15. 154
	NT2RM4001757	38.117	23.659	28. 972	12.593	10.724	21.161	24. 761	19.803
	NY2RM4001758	24. 391	23.518	27.924	5. 579	12.781	14. 153	7. 027	6.943
	NT2804001768	51.099	53.221	60.158	17.044	37. 261	58. 428	34, 390	27. 280
35	NT2RM4001775	15.024	11.154	13.303	2.644	9. 512	9. 892	6. 237	4, 050
	NT2814001776	24. 497	20.843	16. 325	5.116	12.075	8.815	13. 233	6.515
	NY2844001783	44. 218	34.754	35. 521	11.654	27.683	28.899	24. 397	19. 284
	NT2RD4001793	75.698	74.949	146.739	24. 426	38.218	21.996	28. 324	24. 241
	HY2RM4001810	25. 287	22.294	22.627	8.986	12.014	13.754	22.602	19. 691
	NT2804001813	108.290	15.721	11.311	3.071	4. 660	7.061	9.406	9.278
40	NT2RE4001818	55.110	32.332	35. 827	10.603	18. 181	10.893	25. 538	20. 147
40	NT2RD4001819	221.187	103.477	118.661	33.955	61.689	117. 958	105. 557	45. 891
	NT2RE4001823	31.566	19.207	30.580	9.100	12.589	18.948	23.046	12.498
	NT2RM4001828	33.606	37.243	60.904	39.892	17.528	52. 576	22. 264	20.662
	NT2RM4001835	31.946	48. 485	36.681	12.402	10.874	32.404	26.073	33.367
	NT2RH4001836	68.101	53.948	86.019	25. 292	41.216	44. 492	46.063	49.677
	NT2RH4001841	77.551	75.005	64.963	39.736	29, 180	60.179	38. 346	53. 737
45	NY2RH4001842	41.837	31.217	153, 538	19.696	13.432	18.888	13.674	12.515
	DT2824001843	47, 451	47.021	41, 491	12.355	14.857	30.666	19.358	23.477
	RT2RM4001856	35. 284	17.427	22. 905	18.860	0.000	35.066	18.473	17.632
	NT 2 RD 400 1858	34.556	13.809	35. 731	11.606	5, 891	13.370	14. 536	27.815
	NY2854001861	102.500	55. 955	86.639	33.805	25.003	43.868	45.531	30. 143
	NT2R04001863	41.449	33.911	68. 502	24.321	16.482	31.445	31.424	32.578
50		40.706	38.767	51. 589	19.138	24. 325	53. 955	38,078	30.584
	NY2RM4001865				13.720	22.315	49. 946	39.651	110.541
	NT2RM4001869	87.261	35. 753	43.743		23.082	17.690	23.735	26. 533
	HT2RH4001973	31.012	19.677	42. 836	19.140		217.874	135.056	71. 907
	NY2R::14001876	263.450	78.666	162. 933	35.889	80.574		19.933	16.114
	NT2RM4001880	52.575	35. 308	47. 881	20.693	7.377 28.285	39. 267 40. 932	20.399	40. 632
55	NT2RM4001885	62.625	53, 956	164. 215	33.733	1 20.205	40.332	1 20.333	1 40.004

Table 75

	NT2RH4001889	44.826	54. 188	57.058	17. 324	30.679	30.391	31, 401	27.309
	NT2RH4001894	33, 180	21.032	38.644	10. 368	15.617	23.290	26.653	24.028
					11. 443	18.977	24.084	62.995	21.376
5	NT2RM4001897	55. 973	37.135	42.706			19. 933	39.828	71. 231
	NT2RM4001899	79.426	37.833	50.793	22.892	10.010			34. 298
	NT2RM4001905	71.913	42. 987	131,041	19.900	22.521	28.037	22.888	
	NT2RM4001922	68.361	66.765	167.103	32.535	29.282	32.842	21. 101	29.820
	NT2RM4001930	9.761	18.972	11.870	12, 179	5.722	7.704	2.893	19.882
	NT2RM4001938	13.300	9.323	20.059	5. 226	22.340	8.605	6.836	2.737
10	NT2RM4001940	44. 499	28.342	53. 112	22.045	19.769	35.835	24. 329	24.211
70	NT2RM4001942	71.378	109, 603	137. 250	99.314	68.782	123.550	44. 362	143. 236
	NT2RM4001953	73, 750	67.064	218.754	37.265	39.359	37. 249	28.374	31.774
	NT2RI34001965	27.774	33.648	57. 473	21.916	18.921	11.704	7.776	32. 933
	NY2RM4001966	49. 431	24.684	41.501	12. 421	18. 343	29. 179	21.379	18.604
	NT2RM4001969	28.734	22.964	33.007	12.456	14,747	23. 958	15.690	13.553
	NT2RH4001974	82.202	23.827	35. 591	10.813	20.091	38. 983	35. 402	27.290
15	NT2RD4001979	50.759	32.744	64. 327	26.669	29. 268	32. 957	29. 294	45. 426
	NT2RM4001980	64. 506	28. 217	65. 730	29. 832	30.129	51.434	39.037	38. 269
	NT2RM4001984	8. 940	10. 121	18.976	9. 204	7.020	7. 587	10.490	17.931
			27. 2194	64.310	10.713	13.598	56.046	41.155	21.341
	NT2RM4001987	76.782			9.513	9. 423	13.449	15. 551	64. 982
	NT2RM4002013	19.064	9. 935	20. 167	4. 482	9. 866	15. 203	14, 895	11.409
20	NT2RM4002018	23. 330	15. 361	28. 649		36.068	40. 994	22.684	32.604
	NT2RM4002033	103. 629	76.058	255.894	33. 739 25. 591	40. 356	66. 335	30.838	29.885
	NT2RM4002034	97. 025	74.014	204, 281	56, 682	49. 448	68. 685	42.993	58.693
	NT2RM4002044	128. 284	97.260	283. 326		19.793	15, 043	24. 593	16.651
	NT2RU4002047	42.016	31.010	47.604	17. 496	20. 426	36.508	26.858	12.455
	NT2RD4002054	75. 334	24. 437	33.919	5. 362	21.073	24. 192	30. 052	56.881
25	NY2RD4002055	28. 223	41.574	41.231	17.667		26. 659	17.375	42.684
23	HY2RB4002059	24.790	47.792	30,688	32.255	11.889		9.654	12.890
	NT2RM4002061	15. 353	22.159	24.342	33. 358	8.569	13.680	11.468	12.877
	NT2RM4002062	35.603	17.782	25.712	9. 437	13.693	23.679	45, 190	17.589
	NT2RM4002063	106. 902	59. 539	161.049	27. 157	37. 323	44.770	23.625	20. 028
	NT2HM4002066	69. 187	29. 278	44. 089	14. 142	12.777	29.901	19. 168	38. 472
	NT2REM002067	72.915	65.950	164. 446	33.322	23. 243		13. 427	15. 358
30	DT2RE4002073	26.509	19.553	24.129	7.501	12.225	19.453	19.035	9.923
	NY2RY4002074	23.768	16.727	27.356	9.430	10. 288	9. 267	8. 913	5.764
	NT2R34002075	14.729	8.566	14.082	6.113	8. 179	19.921	21.957	5.088
	NT2R14002076	33.772	34, 570	24. 768	12.754	12.370		38. 136	27.441
	NT2RM4002078	65. 837	45. 074	59.931	29.244	28.319	38.890	32.277	29. 982
	NT2R14002081	72.328	49. 374	162.917	29.519	33.925	46.864		4.512
35	NT2RM4002082	31.523	20.963	24. 293	4, 626	7.828	18.917	11.824	15.942
	HT2RM4002093	13.703	12.906	28. 190	14.073	16. 132	8.993		24.718
	NT2RM4002109	48. 477	33,601	44. 587	16. 373	19.020	42.752	31.367	
	HT2RM4002115	52.087	16. 294	25.726	5.046	11.691	15.294	19.312	5.666 8.612
	NT2RM4002118	6.461	10. 205	16.364	2.841	6. 221	5. 928	9. 423	18.787
	NT2RM4002128	24.014	12.586	38.670	8.609	8.704	17.808	16.887 27.756	30. 682
40	NT2RM4002137	60.650	30.735	54.930	9.746	20.827	30.629	18. 198	31.925
	NT2RM4002139	59.820	72. 323	217.660	35.299	32.433	22.926	28. 930	19.620
	HYZRM4002140	61. 939	27. 988	54.095	19.817	18, 951	36.147	54.028	17.830
	HT2RM4002145	55. 935	18.752	37.184	6.758	24. 220	25.455	3, 969	22.927
	NY2RM4002146	10.714	7.232	14.881	2.330	4.463	6.475	8. 266	7.504
	HT2RM4002161	21. 929	10.374	17.604	4. 124	7.983	12.456	10.000	10 755
45	HT2RH4002174	36. 217	21.020	78.760	11.488	14. 155	12.056	10.913	19.766
, 0	HT2RH4002178	51, 201	34. 975		25. 841	26.852	32.083	18.490	38.988
	HT2R14002180	88. 245	86. 565		36.530	47.240	50. 257	25. 291	41.037
	HT2RM4002185	60.374	34.725	47.531	10.870	17.954	36.151	35. 104	14. 336
	NT2R:14002189	443.685	125.746	233.812	62.020	80.189	317.532	213.671	55.692
	NT2RU4002194	110.410	60.176	66.781	14. 262	24. 395	63.199	46, 341	16.491
50	NT2RU4002198	19.112	25. 320		6.006	16.046	10.695	12.653	18.533
50	HY2RM4002205	86.368	52. 183		37.437	37. 350	41.233	35.023	46.891
	HYZRM4002213	87.023	29.632		22. 287	36.169	49.771	58.648	47. 205
	HT2RM4002216	28.034	36.860		61.988	14.040		28.018	31.505
	NT2RM4002226	59, 214	25.842		19.726	22.840		21.306	34. 363
	NY2RM4002237	84.115	47.301		13.185	17.445		282.813	
	NT2RH4002240	21, 140	20.818			4.270	17.260	9. 804	24.400
55	THE PROPERTY	1 2140							

Table 76

	NT2R04002251	39. 895	25.621	38.004	9.808	12.483	27.050	27.880	15.570
	NT2RU4002256	62.880	50. 437	132.459	16.059	20.051	22.911	18. 973	36.148
5	NT2RU4002262	40, 381	19. 221	18.726	4.067	10.643	11.552	18. 506	11.180
5	NT2RM4002266	33. 927	16.247	29.395	7. 271	10.706	15.907	16,746	45.558
	NT2RM4002276	31.555	29. 432	34.470	12. 227	15. 207	18.832	24. 174	41.738
	NT2RM4002278	24, 493	44. 932	54. 554	19. 947	24.631	19.085	14, 211	28. 361
	NT2R04002281	73.045	68.535	120.767	28. 971	77.810	35.833	33. 197	34.350
	NY28M4002287	95. 529	67.191	148. 977	16. 383	32.822	42.647	36. 149	22.550
	NT2RD4002294	37. 325	40.622	32.626	7.879	22. 188	17.681	21.208	18.691
10	NT2R14002298	15. 253	25.056	14. 186	6. 186	12. 213	8.996	13. 334	20. 467
	NT2RD4002301	25. 506	22.524	24. 351	8.779	13.463	11.537	16.605	21.093
	NT2RM4002306	64. 514	27. 130	40.307	8. 697	16.098	30.071	33. 558	17.520
	NT2RM4002323	46. 276	37. 334	108.848	13. 787	18.840	15.998	23.739	23.002
	NT2RM4002334	84, 665	44. 953	240.849	13.009	61.866	67.867	63. 381	16.555
	NT2RM4002339	40. 226	15.664	17.738	4. 286	11.781	13.743	14. 276	7.602
15	NT2RD4002344	15. 209	14, 735	15.127	5. 186	14.835	5.571	6.021	.15.852
	NY2RM4002345	29. 537	16.084	44.040	7. 161	49.725	20.214	15. 169	93.476
	NT2RM4002352	25. 146	26. 320	39.068	10.070	10.828	17.765	20. 622	16.556
	NT2RM4002362	22.727	18. 9671	35.121	7.780	16. 102	13.358	9. 862	21.089
	NY2RM4002373	49.413	25, 049	39.501	16. 293	10.820	16.723	21. 117	10.960
20	NT2RM4002374	45. 312	17.702	80.866	14. 495	13.876	25. 509	12.233	16.564
20	NT2RM4002376	44.035	32.785	33.965	15. 793	15.635	33. 518	17. 499	20.037
	NY2RM4002383	143.921	114.177	338.801	56. 564	36. 130	62. 968	25.071	60.431
	NT 2RM4002390	19. 946	15.647	23. 593	13. 554	0.000	15.764	10.120	21.189
	HT 2RM4002398	33. 574	85,078	55. 577	19.871	29, 143	36.917	34.014	15.071
	NT2RM4002409	62.430	25.690	44, 155	15.629	15. 274	43. 916	36.612	24.609
25	NY2RM4002414	122.797	27.569	49.085	13.732	29. 300	20.609	24. 789	22. 958
20	NT2RE34002438	60. 880	24.210	57. 361	13. 303	21.819	19. 128	27.861	33.288 86.320
	NT2RM4002440	50.958	29.949	58.790	16.516	17.087	22.853	27. 261 43. 072	34.360
	NY28124002446	85. 102	43.893	64.557	15. 166	30.454	59.828 56.095	6. 391	48.088
	NY2814002450	29.806	50.782	20.662	10.226	5.031	21.974	28. 908	14.192
	NT2RE4002452	38. 119	24.046	27.781	13.792	21.980	25. 587	22.709	26.372
30	NT2RM4002457	56.998	45. 958 9. 159	72.065 12.416	21.106 3.859	12.704	4. 423	1.634	7.476
	NT2RM4002458 NT2RM4002450	17. 499 37. 183	7, 502	15.263	2.616	9. 265	20.827	12.805	1.464
	NY 2RM4002464	12.680	10.529	5.512	5. 737	10.707	1.669	5. 391	12.187
	NT2RE4002479	85.068	45. 694	66.175	35. 340	44.661	52. 236	42.316	33.845
	NY 2RM 4002482	714.577	349, 138	482.476	135.984	180.855	462.386	321.086	260.860
	NT2RM4002489	41. 987	36. 475	28. 303	18. 347	20.193	45. 527	22.970	15.427
35	NT2RM4002493	101.547	19.009	34.214	7.129	20.617	58, 926	20.613	6.136
	NY2RM4002499	104.508	114. 364	295. 841	132.961	45.496	125. 546	54.809	138.353
	NT2RM4002504	130.575	85. 186	319.621	58.095	51.615	65. 385	43. 397	39.625
	NT 2RM4002506	17. 534	7.716	22.097	8. 307	8.641	11.973	11.217	19.715
	HY2RM4002510	20. 570	20.274	28. 261	7.195	10.108	9.354	16.982	8. 405
	HT2RM4002527	29.097	14.199	26.008	7.215	11.820	15. 320	15.507	11.537
40	NY2RM4002532	119.266	103.485	252.069	38.479	49, 581	51.534	30.506	48.759
	NT2RE34002534	46.720	29. 222	28.381	12.470	17.005	30.785	27. 381	25. 218 39. 348
	NT2RE4002535	150.736	124.425	370.470	71.472	69.884	70. 122 8. 287	11.868	8.546
	NT2RM4002554	46.680	4.678	15.042	2. 434	7.853	32.697	50.330	26.839
	NT2RE4002558	64. 523	30.756	60.861	17.849	28. 435 13. 855	14. 504	15. 952	20.143
	MT2RM002565	26. 150	21.759		7.961	6.533	14.816	11.242	24.778
45	HY2RH4002567	13.750	9.555	16.128	13. 381	25. 113	37.880	40. 593	30.327
	NT2RM4002571	64. 981	32.370 17.415	44. 482	6. 169	9.094	15.081	8.955	11.463
	NY2R14002572 NY2R14002577	21.932 13.390	34. 537	17.827	8. 379	17. 150	9. 208	20.440	135.375
	NT2RM4002583	43.872	21.818	41.335	7. 938	12.820	25.087	16.879	8.165
	NT2RM4002584	48.978	41.874	47. 589	19.263	15. 387	18.002	26.572	29.591
50	NY2RM4002593	43.140	21.408	34.068	14. 481	17.846	27.459	22.581	18.025
50	NY 2RCM002594	53. 494	32.355	54.474	10.039	23.934	38. 188	30. 209	28.918
	NY 2804002604	49.799	31.218	31.584	14. 197	10.658	52. 255	31. 422	27. 262
	HT2RM4002614	18.848	9. 948	15.663	7.767	10. 103	19. 152	15, 480	10.800
	HT2RM4002515	52.378	28.130	31.691	6. 189	16. 589	25. 551	20.412	22. 945
	NY2RM4002623	31,915	15.505		7.046	11.143	28. 155	15. 957	8. 295
55	NY2RM4002634	27. 202	13.607	23.468	4.566	6.856	27.565	17.040	9. 308
55			<u> </u>						

Table 77

	NT2RH4002636	2.342	5. 234	9.517	3.874	1.465	2.585	2.436	4.543
	NT2RP1000002	114.491	47.508	61.586	25.000	29.448	84.026	73.878	59.624
5	NT2RP1000006	71.057	28.511	44.224	10.202	17.523	40.858	37.373	15. 237
J	NT2RP1000015	7.192	9. 953	16.089	4.506	3.649	7.738	3.651	8.661
	NT2RP1000018	5.882	0.000	0.000	0.000	0.000	2.690	4.737	0.000
	MT2RP1000034	273.802	61.801	59.676	50.413	101.761	283. 598	21.883	51.696
	NT2RP1000035	14.407	14. 328	5.278	5. 331	3.905	19. 347	5. 560	9. 946
	NY2RP1000040	2.229	2.143	2.569	1. 482	0.842	0.251	1.226	0.963
10	NT2RP1000042	2.962	1.516	2.106	0.450	3.003	1.458	1.788	0.000
10	NT2RP1000048	3.312	5. 643	4.404	1,520	1.452	2.742	0.779	17.389
	MT2RP1000050	37.260	7. 381	21.735	7. 969	7.544	14. 598	18.930	19.749
	NT2RP1000056	2.575	8.244	12.209	2.506	2, 248	50.055	1.919	18.856
	NT2RP1000058	7.701	2.152	6.853	1.889	5.740	5.703	5.884	4. 654
	NT2RP1000063	17.863	6.661	8, 488	2.745	0.000	7.494	5. 484	2.401
15	NT2RP1000068	4.612	5. 197	4.140	0.833	1.697	1.068	0.863	1.468
13	NT2RP1000072	143.838	99.413	72.321	37.376	27. 104	99.463	69.787	134.954
	NT 2RP 1000073	1.552	1.742	0.000	0.919	0.996	0.623	4.055	9. 765
	NT2RP1000078	2.896	0.000	0.000	0.230	0.741	0.763	0.567	3. 421
	NT2RP1000079	49.027	29.657	15.514	6.677	6.650	9. 256	18. 182	28. 375
	NT2RP1000080	16.385	13.693	8.875	4. 934	5.832	9.673	15.737	12.194
20	MT2RP1000086	7.169	3.761	10.248	2.945	7.423	5. 286	3.826	0.000
	NT2RP1000087	0.000	5.038	0.000	1.221	3.506	2.887	0.000	2.053
	NT2RP1000089	4. 302	9.012	-8.097	5.674	2.992	4. 524	0.418	13.867 -
	NT2RP1000090	52.428	58.867	69.998	38, 821	17.374	29.637	36.043	79. 235
	MT2RP1000100	3. 207	3.774	1.540	2. 138	1.112	1.149 36.104	0.000 55.994	1.791 56.718
	NT2RP1000101	92.707	46.496	68. 186	33.782	33.861		11.052	2. 965
25	NT2RP1000111	4. 451	9. 940	6.651	2.623	8. 151 0. 000	2.766 1.727	2.041	2. 374
	NY 2RP1000112 NT 2RP1000124	3. 985 24. 505	3.478 9.928	0.000 6.917	2.480 5.644	2553	12.703	2.802	42.544
	NY2RP1000124	24.817	79. 995	139. 555	49.819	97.770	62.060	44. 484	52. 427
	NT2RP1000129	28.170	30.324	26.037	10.799	3.638	16.350	16.315	13. 950
	NT2RP1000130	5. 381	7.279	14. 556	2.578	10.778	12.987	0.000	20.710
	NT2RP1000154	17.054	18.625	18.032	7.765	17.883	13.855	12.502	19.133
30	MY2RP1000163	18.531	7.739	9.822	4.142	3.589	2.512	6.952	17.030
	MY2RP1000170	14.775	6.603	3.911	1.557	5. 549	3.844	7.224	15.609
	NT2RP1000174	10.066	4.006	4.875	1.601	3.951	1.497	5.060	0.857
	NY2RP1000181	108.209	58. 429	137.843	40.129	31.719	74.897	73. 935	56.201
	NT2RP1000191	9. 285	5.645	5. 460	3.099	6.842	12.624	5.864	2.766
	MT2RP1000202	4.547	3.462	7. 203	6. 298	6. 151	3.022	2.481	4. 122
35	NT2RP1000239	0.000	0.000	4.313	1.852	1.396	1.558	2. 101	1.136
	NY2RP1000243	10.228	5. 330	3.864	1.538	6.834	4. 100	5. 184	5. 579
	MT2RP1000255	6.844	3. 187	2.512	1.848	1.326	2.012	5.711	5.678
	NT2RP1000259	10.073	6.510	10.276	1.573	3.601	8.515	4. 509 1. 763	4. 367 0. 000
•	NT2RP1000261	0.000	0.000	0.000	0.000	1.606 78.334	0.000	129.953	95. 341
	NT2RP1000269	233. 453 504. 212	119.331	130.392 684.003	48.933 191.587	126.841	351.080	221. 963	268. 189
40	MT2RP1000272	130. 317	314.887 52.877	78. 345	38. 313	30. 575	71.136	50.465	37. 296
	NT2RP1000279	103. 540	36.699	55. 522	23. 329	29.320	68.415	50.629	9. 388
	NT2RP1000290	383. 695	214, 173	295. 250	136. 106	105.408	257.258	215. 344	195.667
	NT2RP1000293	139. 263	71.666	91.679	43.735	54. 577	85.003	75. 569	61.144
	MT2RP1000300	219.317	94.497	120.961	62.228	73.747	166.238	105. 443	25. 701
45	MY2RP1000324	205. 212	96.463	109.241	73.482	49.779	120.952	75.697	54.085
40	NT2RP1000325	567. 975	208. 141	235. 225	74.690	106.786	296.190	175. 163	181.979
	NT2RP1000326	114.548	37.978	60.587	21.766	22.713	70.707	48.865	22. 186
	NY2RP1000331	14.215	11.082	12.198	9.945	5. 554	9. 595	5. 409	16. 164
	NT2RP1000333	175. 329	62.474	124.398	35.732	30.723	116.009	80.360	48.737
	NY 2RP 1000336	5.071	3.476	0.000	2.085	1.485	4.216	5. 855	5. 234
50	NT2RP1000347	8.732	4.239	0.000	3.444	2.753	3.942	4.829	
-	HT2RP1000348	9.118	3.224	2.495	2.895	3.816	3.756	4.511	1.450
	NT2RP1000349	6.925	4. 441	0.000	1, 180	2.776	3. 407	3.025	2.512
	NY2RP1000353	26. 257	80.510	62.172	39, 139	13.657	50.445	33. 300	
	NT2RP1000356	25. 146	45.385	82. 299	43.972	13. 987	49. 489	26.724	110.239 87.310
	NT2RP1000357	213.820	128. 901	421.667	86. 179		136. 345	94.747	
55 .	NT2RP1000358	186.987	64.055	108.939	32.778	41.723	110. 904	74.510	67.426

## Table 78

									00 (04)
	NT2RP1000360	297. 314	134.601	191.999	71.819	85.890	202.062	147.810	89. 594
	NT2RP1000363	364.040	212.933	280.442 1	36.437	123.748	247.266	256.906	128. 344
									43. 951
	NT2RP1000376	127.768	49.154	84.631	29. 920	40. 910	71.095	82.258	
	HT2RP1000386	39. 353	145.725	56, 520	52. 245	252. 336	185.039	121.336	65. 534
	NT2RP1000407		0.197	0.000	2. 423	0.000	3.032	2.424	3, 461
	NIZKPIUUU4U1	2, 663							0.000
	NT2RP1000409	0.000	5. 878	. 0.000	0.850	0.000	0.424	0.000	
	NT2RP1000413	7, 153	2.048	2,681	0.000	8. 303	4.015	0.344	0.307
					0.000	0,000	0.000	0.000	0.000
	NT2RP1000416	0.000	0.000	0.034					
	NT2RP1000418	9, 174	4. 984	8.733	3.988	5. 668	7.649	7.116	4. 283
	NT2RP1000420	2, 125	0.924	0,000	0.000	0,000	0.000	0, 147	0.000
						0.000	0.189	1.654	0,000
	NT2RP1000434	0.000	19.791	0.000	0.750				
	NT2RP1000439	134. 853	56.272	115,668	51.887	49. 782	73.229	64.079	15. 355
	NT2RP1000443	58. 432	1.440	0.000	3.540	5, 276	7. 299	4. 900	2, 656
•						1, 187	3, 303	1.052	3, 063
	NT2RP1000447	3.820	2.955	0.800	3. 240				
	NT2RP1000448	3. 888	0.697	0.000	0.778	1.043	0.314	0.856	0.000
	NT2RP1000451	5. 766	4.110	3. 245	4.480	1,272	3, 036	1.022	3. 138
			139. 151		114.073	87.709	243.919	188. 141	160.796
	NT2RP1000458	277. 437							
	NT2RP1000460	216. 381	129.722	192.470	86. 161	96. 273	135. 913	170. 172	91.267
	NT2RP1000465	290. 518	221.955	402.881	192.151	210.010	230. 322	182.401	205.887
				61.862	19. 161	13.854	16.791	11.220	11,713
	NT2RP1000468	29. 203	30.933						
	NYZRP1000470	247.991	94.630	118.548	33.073	62. 185	113.536	101. 037	71. 927
	NT2RP1000477	3. 039	1.894	0.000	0.887	1.635	2.721	1.261	1.757
	NT2RP1000478	2.842	0.655	0.000	0.363	1, 122	0.412	1.375	0.000
							0. 993	2.480	1.941
	NT2RP1000481	5. 676	0.693	1.376	2. 294	1, 991			
	NT2RP1000493	5.004	0.820	0.000	1.070	0.687	1. 252	0.401	0. 344
	NT2RP1000513	183. 214	52.178	133.983	29.869	42.569	122.982	62.701	55. 329
				120.005	32.529	32.275	110.978	93.419	62.294
	NT2RP1000522	183. 947	57.483						
	HT2RP1000533	21.686	8. 198	15.700	5, 816	6. 071	12. 902	9. 030	5, 528
	NT2RP1000544	3.732	10.988	1,704	2.465	2, 581	6. 543	9. 371	6.069
			0.310	0.170	0.000	0.000	0.000	0.000	0.000
	NT2RP1000547	0.300						3. 287	1, 199
	NT2HP1000551	3.715	1.322	3.371	0.657	1.870	1.149		
	NT2RP1000567	18, 148	4. 535	7,630	1.128	0.978	9. 115	8. 337	2.192
	NY28P1000574	2.807	2.740	4.159	0.000	1, 266	2.846	0.662	0.000
						4.066	4.517	1.545	3, 168
	NT2RP1000577	5.767	6.059	6. 234	2.033				
	NT2RP1000579	13. 591	6.812	7.808	2.066	3, 452	4.699	7.020	6.279
	NY2RP1000581	23.446	8.664	15.950	5, 531	6.046	15.075	12.761	9,085
					2.580	5.057	9, 162	5.483	15.975
	KT2RP1000593	6.058	14.376	5.780					
	NT2RP1000504	3.081	4. 126	5.413	5. 134	3.748	4. 785	3.835	2.255
	NT2RP1000609	27, 487	3. 174	10,612	2. 228	3, 986	13. 382	13.762	3.825
	NY2RP1000613	4. 356	2. 265	1,529	1.001	0.000	1.184	2.710	0.767
							7. 985	7.518	6.485
	NT2RP1000622	15.005	7. 496	8.013	1.968	1.752			
	NT2RP1000527	17.344	14.772	22.410	6.441	12.047	16.356	20.729	10.336
	NT2RP1000629	15.718	4. 144	12.352	4.104	4. 312	7.820	11.024	7.693
				52.699	15. 138	14.415	30.508	33.741	18, 936
	NT2RP1000530	65. 249	32.499					14.683	16.200
	HT2RP1000639	43. 900	18. 204	18.020	10. 187	10.606	19.791		
	NT2RP1000640	86, 217	156.971	37.078	60.057	32.726	29. 102	17.026	76.883
	NT2RP1000846	7. 394	16.894	13.629	5. 542	5. 660	7.382	1.582	2.851
				53, 935	11.276	9, 11,9	12.945	10.602	15.936
	NY2RP1000559	25.494	13.979						5.755
	NT2RP1000674	10.820	5. 502	9.633	4. 224	4. 542	3.907	5.942	
	NY2RP1000577	187, 310	76.173	99, 589	25.959	49.679	90.146	95. 230	63.227
			5. 907	7.263	2.229	1. 965	2.520	3.853	6.223
	MT2RP1000679	9. 839					17.736	20.738	18.729
	NY2RP1000688	30.741	21.137	41.993	9. 852	14, 205			
	NT2RP1000689	8. 594	2.814	13.021	1.222	4. 171	7.394	4, 473	3.167
	HT2RP1000695	1.813	3.104	2.068	0.810	0.000	0,000	0.786	0.000
						0.000	0.855	0.000	0.607
	NT2RP1000701	1.280	1.032	0.000	0.000				
	NT2RP1000702	4.112	3. 346	8.473	1.156	1.698	1.616	4.749	0.000
	NT2RP1000713	0. 233	0.022	0.927	0.000	0.000	0.000	0.300	0.000
					45. 581	64.142	102.872	121.431	76.919
	MT2RP1000721	199. 987	95. 449	152.563					
	MT2RP1000730	24.414	16. 302	64.370	4.470	6. 129	18.698	8.948	6. 185
	NT2RP1000733	9, 992	13.894	13, 138	3.593	3.087	6. 945	6.918	10.277
				254.026	65.731	120, 196	211.940	169.539	140.421
	NT2RP1000738	357. 351	171. 924					164. 547	67.465
	NY2RP1000739	261. 372	106.684	146.597	37.731	77.574	193.277		
	MT2RP1000740	60.717	34. 534	37.472	15. 130	15.350	35. 255		
				20, 132	3.376	1.635	3.601	3.265	3.969
	NY2RP1000746	13. 275	9. 551	20.132	3 3.310	1.000	1 3.001	, ,,,,,,,	

Table 79

									00.000
NT2RP1	000750	134.663	52. 958	80.346	28.605	36. 158	71.713	92.250	39.685
NTZRPI	000751	17, 717	44. 325	31.941	32. 295	15. 461	19,059	18.084	64.708
	000767	12.860	6.572	9.057	2.510	3.872	3. 120	5, 111	4.085
NT2RP	000769	27.412	21.636	18.089	7.324	7.758	13. 441	12. 436	7.317
NT2RP	000780	7.664	2. 995	3.269	2,715	2.030	0.000	0.000	0.000
	000782	11,618	23. 259	28.607	5.886	16. 596	14.945	5, 301	7,061
								73. 407	26.885
	000796	118. 585	56. 532	75.809	15.096	41.498	78. 341		
NTŽŘP:	000797	215. 680	107. 927	100.844	28.806	53.841	131.952	306. 946	77.792
NTODD	00800	5. 249	3.787	2.211	1.617	7.056	3.306	3, 512	2.851
							16.815	24. 570	12, 101
	000825	49.312	22.623	29.009	4. 529	15. 271			
NT2RP	000833	.67. 848	23, 702	41.132	9, 260	13.328	25. 255	29. 305	27. 307
NT2RP	000834	21. 157	17. 555	15.686	11, 112	11.392	19, 117	14. 348	17. 998
	000836	12.434	11.272	7.839	3. 196	2.621	7.219	5. 827	5. 382
							41. 395	35.068	27. 483
INTERP	1000837	98.743	40.415	104. 822	21.833	23.029			
NT2RP	1000846	14.775	11. 209	35. 656	4.957	5. 131	7.919	3.229	5. 512
	1000847	27. 431	18. 237	16. 588	10.757	10.320	14, 784	19, 182	10.029
			87.847	128. 937	45.113	51.955	144. 598	108.723	51.968
	1000851	214. 374							
NTZRP	1000855	26. 023	29. 514	67.757	23.663	28. 185	38.015	15.874	11.458
NT2RP	1000850	163.711	61.100	101.078	35, 949	41.953	87.889	80. 204	48. 859
	1000902	24. 271	31.899	49.715	12.862	11.237	22.189	17. 326	17.501
						15.239	68. 242	28. 337	14, 115
	1000903	68.716	24. 490	31.806	9. 135				
	1000905	25. 562	13, 385	22. 530	7.568	3.894	6.452	12.011	13. 929
NT28P	1000915	22.768	32. 599	39. 412	17.920	10.752	16.493	10.059	19. 431
	1000916	36. 356	17.076	24.787	8.241	2.752	29.963	18. 336	4.134
	1000921				8. 324	5.717	15. 896	15. 473	4, 407
		20. 200	10.536	22.363					
	1000943	9. 440	4. 278	14.836	10.665	2.682	3. 445	1.686	1. 791
NT2RP	1000944	65, 067	27.816	40.730	12, 441	18.584	43.858	29. 682	34.740
	1000947	18, 414	12.386	22.697	15, 197	10.849	17.723	9.687	20.200
							20. 625	13. 192	15. 554
	1000954	28. 307	24. 912	28. 425	5. 358	10.337			
INT2RP	1000958	21.987	38.788	40.914	23.030	11.285	20. 525	21, 953	28. 920
BT28F	1000959	84. 562	81.956	164. 902	59, 895	32.501	60. 329	47. 308	77.704
	1000966	104, 461	73.705	101.907	58.853	28. 479	65. 560	39, 891	37. 125
							160.514	104.131	41.698
	1000974	213.892	124. 166	171.079	71.813	73.877			
NT2RP	1000980	16.802	11.080	6. 958	4. 146	7.799	7.626	6.311	2.017
NT28F	1000981	50.385	24, 506	35.067	13.841	17.653	24.416	15. 302	5. 946
	1000988	19.623	11.058	22.064	9,003	7.658	18. 310	15. 545	11.394
							27. 348	21. 988	16.177
	1001002	56.891	33.510	22.993	6.717	20.078			
NT2RP	1001004	23.268	13.134	13. 405	6. 295	5. 883	11.999	12. 399	18.783
NT2H	1001007	29. 127	. 10.102	12. 426	8.003	3, 193	18.313	13. 582	9.737
	1001011	36.507	27. 547	42.002	16.657	13.048	28. 628	24.654	12.589
						16.518	29.607	9,620	52. 526
	1001013	9.942	14.082	54. 179	41.030				
ST2RF	1001014	19.677	17. 977	30.913	10.101	11,200	9. 468	17.655	12.776
	1001020	39.078	9. 107	36.274	3.816	13.500	15. 563	15. 121	4.580
		5309.613	985. 566		284.967			3092.785	808. 260
							93.325	67.199	51.245
	1001027	73.098	53. 184	34.629	18.681	24.296			
	1001031	6.727	3. 944	1.413	2.625	2. 583	4.462	2.652	2.043
HT2RE	1001033	34. 383	18. 547	52.827	11.061	12.794	15.798	10.825	16.802
	1001042	16.664	10.042	32.855	18.106	26.513	10. 262	8.945	11.915
	1001045		33.846	51.766	24. 186	48.474	72.682	35. 437	30.767
		189.863							5. 520
	1001073	12. 246	10.612	7.850	6.640	5.048	9.855	6. 935	
NT2RI	1001079	91.852	71.311	176.776	25.199	28.090	49, 291	51, 519	16.408
	1001080	36.634	23. 422	19.061	11.316	14.731	18.812	18.139	11.376
				8. 219	2.444	3, 358	9.872	5, 861	3, 904
	1001113	14. 930	5.617					55.003	98.072
	1001159	327.758	59.111	125. 441	72.993	66.677	187.780		
NT2R	1001173	16.780	13. 137	27.175	6.169	17.090	13.269	9. 476	11. 252
	1001176	12.987	10.035	21.336	6.618	14.457	10.468	9.085	4.024
				35.864	7.900	13, 900	29.446	22. 230	7.579
	1001177	47. 481	25.797						
	1001185	90.471	76.839	221.325	28.708	27.738	39.654	27.055	27.069
HY281	1001199	15.790	17.518	27.913	11.849	14.093	14.390	10.829	11.780
NYMO	1001205	22.415	19.355	38.756	18.438	19.648	28.439	20.497	36.255
							25, 483	15. 692	15, 808
	1001215	26.469	21.856	25.048	13.068	11.039			
NT2R	1001225	54. 629	20. 260	37.472	13.542	10.291	26. 429	33.484	22.194
	1001245	11.787	8. 531	12.195	4.229	4.219	12.906	5.042	9. 166
						1.022	2.368	3. 598	2.028
NT2R	1001247	6, 228	6, 100	7.648	1.747	1.022	2.300	3.030	2.000
		_							

## Table 80

	NT2RP1001248	49. 226	25. 943	116.648	10.461	11.820	12.652	13.256	17.837
		16.172			5.712	7.057	20, 880	11.966	15.830
	NT2RP1001253		14.468	19.494					
5	NT2RP1001286	31.909	17.523	37. 293	9.003	10.973	24. 180	18.180	18.610
3									
	NT2RP1001294	25.024	26.137.1	24.014	7.577	12.732	16. 248	11.737	14.676
	NT2RP1001302	20.570	17.865	14.990	7.914	7.089	11.711	10.424	6.370
	NT2RP1001310	73.669	50.596	61.003	20.191	35. 975	42.746	31.795	30.891
	NT2RP1001311	107.757	35.881	46.474	17.712	21.645	48. 944	43.729	26.945
	NT2RP1001313	55. 324	32.674	63.966	13.492	14.367	18. 129	17.116	14.648
						11.963	16.113	15.675	21.371
10	NT2RP1001324	35. 171	18. 577	22.653	7.819			13.013	
	NT2RP1001349	44. 453	17. 959	25. 475	6.766	11.881	22.818	27.028	20.116
								52.605	27.148
	NT2RP1001361	55. 753	27. 902	58. 131	21.682	28.045	60.728		
	NT2RP1001379	126.769	137.614	71.862	24.018	47,600	154.003	231.914	35.839
								34. 307	
	NT2RP1001385	74. 494	89. 642	123.622	19.403	22.929	45. 989		19.045
	NT2RP1001395	45. 302	31.340	24. 575	7.512	17.756	24. 165	18.832	16.437
15	HT2RP1001410	23.514	23.629	40.104	12.632	9.318	21.843	13.537	8. 295
13	NT2RP1001424	10.618	33.112	10.799	2.636	4.204	7.482	8.833	-25.347
	NY2RP1001432	12.466	40. 995	9.503	1.789	6.323	5. 098	8.187	7.252
	NT2RP1001449	55. 536	20.728	66.767	10.440	26, 188	27. 184	29.004	30.274
	NY2RP1001457	30. 322	32. 7211	37.777	8.330	12.956	20. 340	25. 841	17.849
	NT2RP1001459	88.712	62.417	75.498	27.541	35.602	62.144	51.183	51.852
	NT2RP1001466	16.844	23.355	27.785	10.621	12.274	14. 384	8.050	13.792
20		89.839	111.813	276.258	35.857	23.078	34.083	16.906	15.713
	NT2RP1001475								
	NT2RP1001482	9.804	7. 238	3. 123	7.419	2.367	3. 451	2. 538	1.692
				15, 730	1.433	3.642	8.911	7, 609	6.956
	MY2RP1001494	18. 452	17.405						
	NY2RP1001500	2.143	2.316	3. 634	2.456	0.000	0.086	0.162	0.765
		14.740		16.801	3.704	5. 628	8. 123	9.615	6.297
	NY2RP1001517		13.801						
	NY2RP1001540	50. 226	35.070	52. 423	11.150	17.869	36.090	28. 195	7.025
25						25. 264	48. 623	28.462	18.547
20	NT2RP1001543	87.779	27.665	55.068	12.390				
	NY2RP1001546	51.476	99.385	143.880	25.320	72,799	104. 259	38. 212	42.007
							40.950	26, 433	16.133
	MY2RP1001550	67.741	63. 428	53.684	15.107	31.309			
	NT2RP1001553	34. 956	17.566	22, 966	10.039	10.915	17.367	20.710	19.945
						21.523	42. 121	29.401	24.807
	MT2RP1001555	33. 240	52.576	54.908	25.406	21,323			
	MY 2RP1001563	30. 536	23. 522	26.745	10.623	16.136	20. 228	17.699	11.340
00									
30	NT2RP1001569	90.271	31.802	37.662	7.791	18.755	32. 159	31.572	22.545
	NT2RP1001584	125.503	64.642	101.860	20.979	38.153	69. 983	85.177	68.021
	NY2RP1001599	25. 536	22.635	29.822	7.141	9.376	19.848	14.150	13.608
	NY2RP1001616	38.077	18. 321	20. 981	7.268	5, 256	12.873	14.067	12.210
	NT2RP1001654	77.215	24. 275	26.850	14.308	14.684	36.754	26.803	17.786
	NY2RP1001665	20, 132	15, 451	16.433	5. 156	9.958	5.979	8.761	8.109
35	NT2RP1001679	251.384	264.730	245. 821	192. 156	85.798	197.731	172.668	434.739
	NY2RP1001681	21.960	21.892	16.974	17. 231	5. 379	21.608	10.982	20.811
	NT2RP1001694	27.832	32. 368	36.517	12.438	29, 150	109.147	231.086	69.267
	NY2RP2000001	79.348	34. 825	26.858	8.546	17.604	24. 165	27.629	18.039
	MT2RP2000006	32.218	26.701	47.407	11.066	8.723	14. 994	13. 215	12.652
	NT2RP2000007	54. 262	32.503	34.116	12.829	11.972	20, 410	21.705	11.281
40	NT2RP2000008	34.810	31.036	59.562	20.809	17. 226	20.509	17. 286	54.391
	NY 2RP2000010	12.320	9.820	24.557	3.019	5.341	8.149	10.076	5.865
	TOTAL LOCALIO						64. 567	50. 108	46.745
	NT2RP2000011	121.718	115. 419	216.553	41.153	44.035	04. 30/		
	NT2RP2000027	74.085	69.757	136.369	23.981	28. 217	40.308	24. 108	20.710
	NT2RP2000028	23.699	28. 386	27.077	10.607	11.433	22.532	14. 265	11.554
	NT2RP2000032	10.199	6.568	16.529	6.282	6.462	9.523	8, 119	8.527
									1
45	NT2RP2000040	383.423	222.501	199.099	79.455	81.787	229.220	181.239	162.128
45	NT2RP2000042	97,011	62.254	67.677	29.525	13.003	45. 921	45.196	41, 158
	MT2RP2000045	73.700	49.722	66.899	21.221	17. 180	32.492	32.785	35. 403
	NT28P2000051	37, 323	46.342	93. 958	33. 924	13.292	43.534	29. 174	17. 962
	HT2RP2000054	99.806	54.072	69.945	21.897	22.707	40.001	40.807	38.782
	NY2RP2000056	57.518	40.207	41.868	18. 309	24. 303	26.794	25.564	25. 156
	NT2RP2000057	156.050	177.739	178.741	136.241	76.886	130.744	163. 333	207. 593
50	NT2RP2000087	59, 366	13.414	39.371	6. 372	16.511	22.699	22.699	5.023
	NT2RP2000070	107.618	50. 674	57.709	17.458	29.909	83.478	48.688	26.235
	NT2RP2000076	48.409	27. 260	29.570	12.733	8. 235	32.852	11.701	13.485
								-	
	NT2RP2000077	94. 993	53. 327	77.668	25.110	14.024	49, 100	33.647	31, 168
	NT2RP2000079	62.685	66.203	139. 230	32.930	26.739	30.432	16.329	18.678
	NY2RP2000088	71.164	29, 601	52.899	11.567	20.381	42.871	35. 756	8.836
55				•					
JJ									

# Table 81

					45 466	17 150	74 252 1	10 714 1	15 027 T	10, 202
5										
No.										
HITEPPRODUIDS   156, 612   136, 647   138, 5078   30, 234   79, 342   81, 573   54, 191   92, 488   HITEPPRODUID   22, 648   27, 255   23, 255   8, 385   6, 127   16, 427   11, 222   29, 326   187287900116   32, 747   25, 306   35, 305   21, 085   8, 128   21, 812   11, 292   29, 326   31, 327   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 327   32, 321   31, 321   32, 324   31, 358   31, 335   35, 347   39, 360   31, 365	5									
1172#P7000110   12, 814   21, 255   23, 551   8, 385   5, 127   16, 427   11, 227   18, 744     1172#P7000118   22, 474   72, 306   33, 530   2, 508   3, 128   21, 812   11, 122   29, 258     1172#P7000119   28, 773   75, 708   213, 188   30, 879   26, 575   32, 244   18, 663   22, 213     1173#P7000120   28, 184   40, 4072   11, 425   17, 144   20, 374   18, 788   14, 212     1173#P7000128   68, 255   51, 174   75, 174   25, 719   22, 146   30, 674   19, 806   13, 046     1173#P7000138   68, 255   51, 174   75, 174   25, 719   22, 146   30, 674   19, 806   13, 046     1173#P7000137   72, 104   61, 190   75, 784   21, 435   31, 539   75, 147   46, 430   37, 718     1173#P7000137   72, 104   61, 190   75, 784   21, 435   31, 539   75, 147   46, 430   37, 718     1173#P7000137   72, 136   61, 190   75, 784   21, 435   31, 539   75, 147   46, 430   37, 718     1173#P7000137   72, 136   81, 737   200, 882   37, 908   35, 422   38, 443   88, 450   20, 252     1173#P7000138   11, 13, 14, 146   13, 146   14, 145   14, 145   12, 284   22, 086   12, 19   10, 751     1173#P7000139   72, 131   11, 646   139, 508   3, 97   5, 183   6, 465   8, 554   32, 06     1173#P7000131   35, 435   45, 147   17, 198   15, 404   30, 144   34, 404   30, 404   3										
TETERPRODOTIS   2x, 2xt   7x, 300   35, 305   21, 085   3, 128   21, 812   11, 292   29, 328   117287200178   37, 77, 780   213, 188   30, 197   26, 975   32, 244   18, 788   31, 212   11, 212										
10										
10		MT2RP2000116								
Time		MT2RP2000119	87.773							
ST2RP2000133	10	MT2RP2000120		40. 341						
HTTSPP20001747   127. 104   61. 1807   75. 784   21. 438   33. 839   75. 147   46. 430   37. 71. 81	70	NT2RP2000126		51.174	75.714					
TYPEPTOOD156		MT2RP2000133	40.974	21.406						
NTTRP7000156		MT2RP2000147	121.104	61.190	75.784		33.839			
15 NTTRP2000161 9.493 12.579 24.5715 5.678 7.14.121 12.284 22.086 12.179 10.763 NTTRP20001715 7.83 12.579 24.5715 5.678 7.191 9.079 10.000 10.000 NTTRP2000175 78.383 12.579 24.5715 5.678 7.191 9.079 9.091 65.465 17.000 NTTRP2000175 78.383 44.514 77.095 15.404 30.614 50.131 40.311 40.201 NTTRP2000176 78.383 44.514 77.095 15.404 30.614 50.131 40.311 40.201 NTTRP2000178 60.513 42.174 41.514 14.454 19.558 28.068 22.431 40.205 NTTRP2000183 10.319 90.798 19.000 41.414 41.454 19.558 28.068 22.431 15.249 NTTRP2000183 10.319 90.798 19.000 41.414 10.656 27.518 41.68 44.54 16.271 60.391 52.249 NTTRP2000204 91.419 106.652 (25.3856 91.981 356.252 154.885 68.553 248.768 NTTRP2000204 91.419 106.652 (25.3856 91.981 356.252 154.885 68.553 248.768 NTTRP2000204 91.419 106.652 (25.3856 91.981 356.252 154.885 68.553 248.768 NTTRP2000204 69.052 62.644 64.951 28.002 14.255 42.146 33.510 11.634 NTTRP2000221 69.052 62.038 16.519 19.712 16.655 28.186 30.382 15.64 NTTRP2000221 69.052 62.038 16.519 19.712 16.655 28.186 30.382 15.64 NTTRP2000221 22.7466 16.377 180.416 68.560 85.769 155.055 119.086 87.184 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 43.941 52.603 20.024 20.149 47.211 52.894 32.775 NTTRP2000231 74.158 13.393 33.88 42.776 8.787 NTTRP2000231 74.158 13.393 33.88 42.776 8.787 NTTRP2000231 74.158 13.393 33.88 42.776 NTTRP2000231 74.158 13.393 33.89 13.797 14.797 15.654 21.302 17.214 20.486 NTTRP2000231 74.158 13.393 33.89 12.797 12.706 9.712 52.991 13.775 NTTRP2000231 74.158 13.393 33.893 13.494 13.493 13.493 13.493 13.294 13.595 13.294 13.595 13.294 13.795 NTTRP2000231 74.588 33.593 13.498 13.494 13.595 13.393 13.22 46.624 13.795 NTTRP2000231 74.589 33.595 13.654 13.393 13.22 46.624 13.795 NTTRP2000231 74.589 13.59		NT2RP2000153	96. 598	63.476	66.144	23.377				
15		NT2RP2000156	115, 309	87.737	200. 582	37.008				
No.		NT2RP2000157	24. 318	18.096	28.697	14. 121	12.284			
STERP2000173   228, 420   98, 033   150, 036   37, 188   58, 850   14, 315   90, 491   66, 485   817RP2000175   78, 339   44, 514   77, 1036   15, 404   30, 614   50, 131   40, 431   4	15	NY2RP2000161		12.679	24. 575	5.678	7. 191		8. 105	4
20   ST2RP2000178   T3, 339   44, 514   71, 095   15, 404   30, 614   50, 131   40, 431   40, 206   ST2RP2000188   120, 139   30, 788   139, 074   34, 168   44, 541   64, 271   60, 391   53, 828   ST2RP2000189   51, 304   70, 037   704, 874   30, 805   27, 733   45, 334   28, 749   18, 697   ST2RP20002005   30, 577   27, 665   61, 321   18, 312   9, 596   77, 099   7, 227   6, 812   ST2RP20002005   30, 577   27, 665   61, 321   18, 312   9, 596   77, 099   7, 227   6, 812   ST2RP20002005   30, 577   27, 665   61, 321   18, 312   9, 596   77, 099   7, 227   6, 812   ST2RP20002005   53, 204   84, 346   85, 469   22, 464   20, 371   37, 091   31, 135   31, 123   ST2RP2000220   65, 520   38, 161   51, 891   19, 712   16, 665   28, 186   80, 382   25, 164   ST2RP2000230   53, 204   84, 346   85, 469   22, 464   20, 371   37, 091   31, 135   31, 123   ST2RP2000231   237, 426   116, 377   180, 416   88, 560   85, 769   155, 655   119, 686   87, 184   772P2600231   237, 426   116, 377   180, 416   88, 560   85, 769   155, 655   119, 686   87, 184   772P2600233   32, 386   15, 393   30, 197   8, 574   4, 025   17, 013   20, 268   32, 735   872P2600233   37, 158   43, 941   52, 503   20, 024   20, 149   47, 211   52, 894   52, 735   872P2600233   37, 158   43, 941   52, 503   20, 024   20, 149   47, 211   52, 894   52, 735   872P2600234   47, 308   13, 339   13, 368   4, 823   12, 697   8, 493   12, 292   8, 218   372P2600234   47, 308   31, 339   13, 368   4, 823   12, 697   8, 498   31, 322   46, 624   872P2600254   47, 308   31, 339   31, 368   4, 823   12, 697   8, 498   31, 322   46, 624   872P2600254   44, 605		NT2RP2000168	11, 413	14.646	19.908	3. 979	5. 383	6.466	8.554	
NTZRPZ000188   60. 513   42. 174   41. 514   14. 454   19. 558   28. 068   22. 439   15. 249		NT2RP2000173	228. 420	98.033	150.036	37. 188	58.850	114.315	90. 491	
SITZRP200018				44.514		15.404	30.614	50.131	40. 431	40. 206
STZRPZOBOT98				42.174	41.614	14. 454	19.558	28. 068	22.439	16.249
ST2RP2000205			120, 139	90.798	139.074	34. 168	44.541	64. 271	60.391	
STZRPZ0002205   30, 577   27, 66.5   61, 321   18, 312   9, 596   17, 099   7, 227   6, 812	20		91.304	70.037	204.874	30.805	27.133			
## ## ## ## ## ## ## ## ## ## ## ## ##		NT2RP2000204	91.419	106.652	263.856	91.981	356.822	154.895		
STZRPZ000224   69.062   62.644   64.951   28.002   14.255   42.146   33.510   51.634     STZRPZ000230   56.320   38.161   51.891   19.712   16.865   28.186   30.382   25.184     STZRPZ000231   237.426   116.377   160.416   68.560   85.769   155.055   119.086   87.184     STZRPZ000232   49.708   32.849   24.700   10.366   11.881   31.935   21.623   13.775     STZRPZ000239   32.380   15.399   30.197   8.574   4.025   17.013   20.268   23.735     STZRPZ000239   32.380   15.399   30.197   8.574   4.025   17.013   20.268   23.735     STZRPZ000230   49.173   38.363   78.202   15.737   15.654   21.302   17.214   20.486     STZRPZ000230   37.308   13.339   13.368   4.823   12.687   8.493   12.992   9.218     STZRPZ000256   37.550   25.977   25.477   12.706   9.212   23.055   14.601   18.126     STZRPZ000258   39.114   41.740   49.525   15.968   19.509   29.341   17.580   17.049     STZRPZ000258   39.114   41.740   49.525   15.968   19.509   29.341   17.580   17.049     STZRPZ000259   39.114   41.740   49.525   15.968   19.509   29.341   17.580   17.049     STZRPZ000274   15.514   7.310   20.284   4.327   6.428   13.479   7.807   4.833     STZRPZ000277   12.320   12.198   8.692   2.395   5.014   49.489   25.505   41.022     STZRPZ000279   12.294   6.735   9.825   2.486   5.467   4.265   7.545   6.898     STZRPZ000279   38.289   22.877   35.809   11.594   41.150   24.652   25.978   24.657     STZRPZ000279   76.236   77.277   20.685   45.399   40.991   22.703   77.905     STZRPZ000279   76.236   77.277   20.685   45.399   43.91   44.507   27.804     STZRPZ000279   76.236   77.277   20.685   45.399   43.90   40.991   22.703   77.905     STZRPZ000289   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     STZRPZ000289   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     STZRPZ000289   51.997   39.585   39.607   47.746   39.914   36.153   31.476   28.603     STZRPZ000337   45.414   46.20   50.933   46.420   15.624   43.030   26.358   27.011   31.395     STZRPZ000337   37.498   20.		NT2RP2000205	30. 577	27.665	61.321	18.312	9.596	17.099	7.227	
STZRP2000230   55. 320   38. 161   51. 891   19. 712   16. 865   28. 186   30. 382   25. 164     STZRP2000231   237. 426   116. 377   160. 416   68. 560   85. 769   155. 055   119. 086   87. 184     STZRP2000233   74. 158   43. 941   52. 603   20. 024   20. 149   47. 211   52. 894   52. 273     STZRP2000239   32. 380   15. 399   30. 197   85. 574   4. 022   17. 013   20. 268   23. 735     STZRP2000230   44. 173   38. 363   78. 202   15. 737   15. 654   21. 302   17. 214   20. 486     STZRP2000236   37. 530   25. 977   25. 477   17. 766   9. 212   23. 055   44. 601   18. 126     STZRP2000236   37. 650   25. 977   25. 477   17. 766   9. 212   23. 055   44. 601   18. 126     STZRP2000257   69. 335   66. 181   244. 979   45. 881   37. 192   46. 969   31. 322   46. 624     STZRP2000253   39. 114   41. 740   49. 525   15. 968   19. 509   29. 341   17. 580   17. 049     STZRP2000254   46. 051   30. 214   48. 737   10. 438   13. 441   22. 574   19. 894   19. 558     STZRP2000257   73. 075   55. 962   155. 102   33. 557   26. 014   49. 469   25. 505   41. 022     STZRP2000277   73. 075   55. 962   155. 102   33. 557   26. 014   49. 469   25. 505   41. 022     STZRP2000277   12. 320   12. 198   8. 692   2. 395   5. 097   7. 436   9. 834   3. 452     STZRP2000279   12. 294   6. 735   9. 825   2. 486   5. 467   4. 265   7. 545   6. 88     STZRP2000279   12. 294   6. 735   9. 825   2. 486   5. 467   4. 265   7. 545   6. 88     STZRP2000279   12. 294   6. 735   9. 825   2. 486   5. 467   4. 265   7. 545   6. 88     STZRP2000279   12. 294   6. 735   9. 825   2. 486   5. 467   4. 265   7. 545   6. 88     STZRP2000279   12. 320   12. 198   8. 692   2. 395   5. 097   7. 436   9. 834   3. 452     STZRP2000279   12. 320   12. 198   8. 692   2. 395   5. 097   7. 436   9. 834   3. 452     STZRP2000327   12. 294   6. 735   9. 825   2. 486   5. 467   4. 265   7. 545   6. 88     STZRP2000328   38. 329   22. 877   38. 89   11. 594   14. 150   24. 632   25. 978   24. 657     STZRP2000328   38. 600   35. 521   50. 933   15. 515   20. 30					85. 459	22.464	20.371	37.407		
STZRP2000230			69,062		64.951		14. 265	42.146	33.510	51.634
STZRPZ000231   237,426   116,377   150,416   68,560   85.769   155.055   119,086   87.184     NTZRPZ000232   49,708   32,849   24,700   10,366   11,881   31,935   21,623   13,775     STZRPZ000233   74,158   43,941   52,503   20,024   20,149   47,211   52,894   52,273     STZRPZ000239   32,380   15,399   30,197   8,574   4,025   17,013   20,268   23,735     STZRPZ000240   49,173   38,363   78,202   15,737   15,654   21,302   17,214   20,486     STZRPZ000256   37,650   25,977   25,477   12,706   9,212   23,055   14,601   18,126     STZRPZ000257   59,335   66,181   244,979   44,881   37,192   46,569   31,322   46,624     STZRPZ000258   39,114   41,740   49,525   15,968   19,509   29,341   77,580   17,049     STZRPZ000274   73,075   55,962   155,102   33,557   26,014   49,469   26,555   41,022     STZRPZ000277   73,075   55,962   155,102   33,557   26,014   49,469   26,555   41,022     STZRPZ000277   13,200   12,198   8,692   2,395   5,097   7,436   9,834   3,452     STZRPZ000278   33,244   49,998   59,636   18,166   19,261   33,588   39,787   48,770     STZRPZ000279   12,294   6,735   9,825   2,486   5,467   4,265   7,545   6,898     STZRPZ000279   12,294   6,735   9,825   2,486   5,467   4,265   7,545   6,898     STZRPZ000279   12,294   6,735   9,825   2,486   5,467   4,265   7,545   6,898     STZRPZ000279   12,294   6,735   9,825   2,486   5,467   4,265   7,545   6,898     STZRPZ000279   12,294   6,735   9,825   1,594   4,1150   24,632   25,978   48,770     STZRPZ000289   51,397   39,352   53,601   14,746   19,914   36,153   11,476   28,603     STZRPZ000297   76,236   71,227   206,854   45,833   34,290   40,991   22,703   77,905     STZRPZ000300   29,075   14,696   16,125   5,503   10,245   16,627   19,121   1,456     STZRPZ000337   31,686   30,551   30,300   31,272   27,377   10,155   35,151   31,360   31,378   31,389   26,630   31,322   27,398   33,493   22,925   32,863     STZRPZ000337   31,894   26,635   31,577   17,745   41,448   31,500   31,48   7,591   12,275   37,981   31,481   31,412   27,988   31,429					51.891	19.712	16.865		30. 382	
NTZRPZ000233	25			116.377	160.416	68, 560	85.769	155.055	119.086	
NTZRPZ000233   32.380   15.399   30.197   8.574   4.025   17.013   20.268   23.735		NT2RP2000232	49.708	32.849	24.700	10.366	11.881			
NTZRPZ000260		NT2RP2000233	74. 158	43.941	52.603	20.024				
		NT2RP2000239	32.380	15.399	30.197	8. 574	4.025			
ST2RP2000258   37.650   25.977   25.477   12.706   9.212   23.055   14.601   18.126     ST2RP2000257   69.335   66.181   244.979   45.881   37.192   46.969   31.322   46.524     ST2RP2000258   39.114   41.740   49.525   15.968   19.509   29.341   17.580   17.049     ST2RP2000270   73.075   55.962   155.102   33.557   26.014   49.469   26.505   41.022     ST2RP2000277   73.075   55.962   155.102   33.557   26.014   49.469   26.505   41.022     ST2RP2000277   12.320   12.198   8.692   2.395   5.097   7.436   9.834   3.452     ST2RP2000279   12.294   6.735   9.825   2.486   5.467   4.265   7.545   6.898     ST2RP2000289   36.324   49.998   59.636   18.166   19.261   33.585   39.787   48.270     ST2RP2000289   36.329   22.877   35.809   11.594   41.150   24.632   25.978   24.657     ST2RP2000289   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     ST2RP2000289   76.236   71.227   206.854   45.839   34.290   40.991   22.703   77.905     ST2RP2000299   76.236   71.227   206.854   45.839   34.290   40.991   22.703   77.905     ST2RP2000390   28.739   29.954   34.444   15.641   0.562   21.620   14.607   21.804     ST2RP2000329   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.416     ST2RP2000329   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.416     ST2RP2000337   30.149   22.042   28.730   11.084   7.733   16.593   11.667   6.892     ST2RP2000386   21.429   15.884   19.766   6.532   11.361   9.148   7.691   12.275     ST2RP2000386   21.429   15.884   19.766   6.532   11.361   9.148   7.691   12.275     ST2RP2000387   20.133   21.4281   190.587   52.114   101.706   157.153   153.536   79.610     ST2RP2000386   21.429   15.884   19.766   6.532   11.361   9.148   7.691   12.275     ST2RP2000386   21.429   15.884   19.766   6.532   11.361   9.148   7.691   12.275     ST2RP2000386   21.429   15.884   19.766   6.532   11.361   9.148   7.691   12.275     ST2RP2000387   20.313   20.422   20.422   20.422   20.422   20.423   20.424   20.424   20.655   20.303   11.		NT2RP2000240	49. 173	38. 363	78.202	15.737	15.654			
NTZRPZ000257   69.335   66.181   244.979   45.881   37.192   46.969   31.322   46.624     NTZRPZ000258   39.114   41.740   49.525   15.968   19.509   29.341   17.580   17.049     NTZRPZ000270   73.075   55.962   155.102   33.557   26.014   49.469   26.505   41.022     NTZRPZ000270   73.075   55.962   155.102   33.557   26.014   49.469   26.505   41.022     NTZRPZ000277   12.320   12.198   8.692   2.395   5.997   7.436   9.834   3.452     NTZRPZ000279   12.320   12.198   8.692   2.395   5.997   7.436   9.834   3.452     NTZRPZ000263   63.324   49.998   59.636   18.166   19.251   33.586   39.787   48.270     NTZRPZ000264   38.289   22.877   35.809   11.594   14.150   24.632   25.978   24.657     NTZRPZ000289   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     NTZRPZ000299   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     NTZRPZ000299   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     NTZRPZ000290   28.739   29.954   34.444   15.641   10.552   21.620   14.607   21.804     NTZRPZ0003127   54.414   66.201   24.879   17.704   13.651   22.202   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   22.222   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   22.222   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   24.922   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   24.922   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   24.922   17.858   30.618     NTZRPZ000327   45.414   66.201   24.879   17.704   13.651   24.922   17.858   30.618     NTZRPZ000327   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.438     NTZRPZ000327   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.438     NTZRPZ000337   14.768   17.723   21.977   6.203   6.291   12.498   8.041   5.971     NTZRPZ000356   82.288   25.117   44.596   10.329   16.344   44.774   37.686   6.290     NTZRPZ0003		NT2RP2000248	17, 308	13.339	13.368		12.687			
NTT2RP2000250	30	NT2RP2000256	37.650	25, 977	25.477		9.212			
NT2RP2000261   46.051   30.214   48.737   10.438   13.441   22.674   19.894   19.556     NT2RP2000270   73.075   55.962   155.102   33.557   26.014   49.469   26.505   41.022     NT2RP2000277   15.514   7.310   20.284   4.327   6.428   13.479   7.807   4.833     NT2RP2000277   12.320   12.198   8.692   2.395   5.097   7.436   9.834   3.452     NT2RP2000279   12.294   6.735   9.825   2.486   5.467   4.265   7.545   6.898     NT2RP2000283   63.324   49.998   59.636   18.165   19.261   33.586   39.787   48.270     NT2RP2000283   38.289   22.877   35.809   11.594   41.150   24.632   25.978   24.557     NT2RP2000285   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     NT2RP2000289   28.739   29.954   34.444   15.641   10.562   21.620   14.607   21.804     NT2RP2000310   29.075   14.696   16.125   5.503   10.245   16.627   19.121   11.456     NT2RP2000328   36.600   35.521   50.933   15.515   23.798   33.981   22.925   32.863     NT2RP2000328   45.820   29.353   14.172   22.985   15.584   34.848   35.626   29.436     NT2RP2000337   45.820   29.353   14.172   22.985   15.584   34.848   35.626   29.436     NT2RP2000337   45.820   29.353   14.172   22.985   15.584   30.638   30.618     NT2RP2000337   45.820   29.353   14.172   22.985   30.638   30.618     NT2RP2000337   45.820   29.353   14.172   22.985   30.638   30.618   31.292   32.863     NT2RP2000337   30.149   22.042   28.730   11.084   7.733   16.593   11.567   6.892     NT2RP2000336   62.288   08.853   14.700   2.291   4.114   11.789   8.150   7.184     NT2RP2000366   82.288   25.117   44.596   10.329   16.344   44.774   37.686   6.295     NT2RP2000386   21.429   15.884   19.0587   52.114   101.706   157.153   153.356   79.610     NT2RP2000386   21.429   15.882   21.557   30.952   32.657   39.766     NT2RP2000386   21.429   15.862   21.685   34.558   45.976   138.158   139.412   79.987     NT2RP2000386   21.429   15.862   21.685   25.253   30.952   32.657   39.766     NT2RP2000386   21.429   15.685   25.553   34.558   45.976   138.158	00	NT2RP2000257	69. 335	66.181						
NTZRPZ000270		NT 2RP2000258	39, 114	41.740		15.968				
NYZRPZOOUZ74		NT2RP2000261	46.051	30.214	48.737	10.438	13.441			
NT2RP2000277   12.320   12.198   8.692   2.395   5.097   7.436   9.834   3.452     NT2RP2000279   12.294   6.735   9.825   2.486   5.467   4.265   7.545   6.898     NT2RP2000283   63.324   49.998   59.636   18.166   19.261   33.586   39.787   48.270     NT2RP2000280   38.289   22.877   35.809   11.594   14.150   24.632   25.978   24.657     NT2RP2000280   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     HY2RP2000297   76.236   71.227   206.854   45.839   34.290   40.991   22.703   77.905     NT2RP2000396   28.739   29.954   34.444   15.641   10.562   21.620   14.607   21.804     NT2RP2000310   29.075   14.696   16.125   5.503   10.245   16.627   19.121   11.456     NT2RP2000327   45.414   16.201   24.879   17.704   13.651   24.922   17.858   30.618     NT2RP2000328   36.600   35.521   50.933   15.515   23.798   33.981   22.925   32.863     NT2RP2000329   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.436     NT2RP2000337   14.768   17.723   21.972   6.203   6.291   12.498   8.041   5.971     NT2RP2000336   55.051   82.391   46.420   15.624   13.030   26.358   27.011   51.395     NT2RP2000356   30.149   22.042   28.730   11.084   7.733   16.593   11.667   6.892     NT2RP2000368   25.288   25.117   44.596   10.329   16.344   44.774   37.686   6.290     NT2RP2000368   21.429   15.884   19.746   6.552   11.361   9.148   7.691   12.275     NT2RP2000369   21.429   15.884   19.746   6.552   11.361   9.148   7.691   12.275     NT2RP2000386   25.303   11.496   120.655   34.558   45.976   138.158   139.412   79.987     NT2RP2000386   23.332   142.481   190.587   52.114   101.706   157.153   153.536   79.610     NT2RP2000394   31.766   23.882   31.577   11.745   14.448   23.860   24.285   20.279     NT2RP2000395   231.332   142.481   190.587   52.114   101.706   157.153   153.536   79.610     NT2RP2000412   67.028   66.250   119.740   21.685   25.553   30.952   32.657   39.766   87.287   88.004   87.787   88.004   87.787   88.004   87.787   88.004   87.787   88.004   87.787		NT 2RP2000270	73.075							
NT2RP2000283   63, 324   49, 998   59, 636   18, 166   19, 261   33, 586   39, 787   48, 270     NT2RP2000283   63, 324   49, 998   59, 636   18, 166   19, 261   33, 586   39, 787   48, 270     NT2RP2000289   51, 997   39, 352   53, 601   1, 594   14, 150   24, 632   25, 978   24, 657     NT2RP2000299   76, 236   71, 227   206, 854   45, 839   34, 290   40, 991   22, 703   77, 905     NT2RP2000290   28, 739   29, 954   34, 444   15, 641   10, 562   21, 620   14, 607   21, 804     NT2RP2000310   29, 075   14, 696   16, 125   5, 503   10, 245   16, 627   19, 121   11, 456     NT2RP2000327   45, 414   16, 201   24, 879   17, 704   13, 651   24, 922   17, 888   30, 618     NT2RP2000328   36, 600   35, 521   50, 933   15, 515   23, 798   33, 981   22, 925   32, 863     NT2RP2000337   45, 820   29, 353   14, 112   22, 985   11, 584   34, 848   35, 626   29, 436     NT2RP2000337   44, 46   46, 46, 47, 47, 47, 47, 47, 47, 47, 47, 47, 47		HY2RP2000274								
NTZRP2000283   38, 289   22, 877   35, 809   11, 594   14, 150   24, 632   25, 978   24, 657	25									
HT2RP2000289 38.285 22.877 35.809 11.594 14.150 24.632 25.978 24.657  NT2RP2000289 51.997 39.352 53.601 14.746 19.914 36.153 31.476 28.603  RT2RP2000297 76.236 71.227 206.854 45.839 34.290 40.991 22.703 77.905  NT2RP2000398 28.739 29.954 34.444 15.641 10.562 21.620 14.607 21.804  HT2RP2000310 29.075 14.696 16.125 5.503 10.245 16.627 19.121 11.456  NT2RP2000327 45.414 16.201 24.879 17.704 13.651 24.922 17.858 30.618  NT2RP2000328 36.600 35.521 50.933 15.515 23.798 33.981 22.925 32.863  NT2RP2000328 45.820 29.353 14.112 22.985 11.584 34.848 35.626 29.436  NT2RP2000333 33.894 26.367 89.382 12.302 13.127 27.377 10.155 15.517  NT2RP2000346 53.051 82.391 46.420 15.624 13.030 25.358 27.011 31.395  NT2RP2000346 53.051 82.391 46.420 15.624 13.030 25.358 27.011 31.395  NT2RP2000368 82.288 25.117 44.596 10.329 16.344 44.774 37.686 6.290  ETZRP2000369 21.429 15.884 19.746 6.532 11.361 9.148 7.691 12.275  NT2RP2000383 31.766 23.882 31.577 11.745 14.448 23.860 24.285 20.279  NT2RP2000386 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610  NT2RP2000386 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610  NT2RP2000341 97.169 86.021 59.155 47.116 24.169 74.619 64.790 62.555  NT2RP2000412 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766  NT2RP2000412 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766	30	NT2RP2000279	12. 294							
NT2RP2000289   51.997   39.352   53.601   14.746   19.914   36.153   31.476   28.603     NY2RP2000297   76.236   71.227   206.854   45.839   34.290   40.991   22.703   77.905     NT2RP2000298   28.739   29.954   34.444   15.641   10.562   21.620   14.607   21.804     NT2RP2000310   29.075   14.696   16.125   5.503   10.245   16.627   19.121   11.456     NT2RP2000327   45.414   16.201   24.879   17.704   13.651   24.922   17.858   30.618     NT2RP2000327   45.414   16.201   24.879   17.704   13.651   24.922   17.858   30.618     NT2RP2000329   45.820   29.353   14.112   22.985   11.584   34.848   35.626   29.436     NT2RP2000333   33.894   26.367   89.382   12.302   13.127   27.377   10.155   15.517     NT2RP2000337   14.768   17.723   21.972   6.203   6.291   12.498   8.041   5.971     NT2RP2000336   53.051   82.391   46.420   15.624   13.030   26.358   27.011   31.395     NT2RP2000357   30.149   22.042   28.730   11.084   7.733   16.593   11.667   6.892     NT2RP2000366   82.288   25.117   44.596   10.329   16.344   44.774   37.686   6.290     NT2RP2000366   82.288   25.117   44.596   10.329   16.344   44.774   37.686   6.290     NT2RP2000376   205.303   11.496   120.655   34.558   45.976   138.158   139.412   79.987     NT2RP2000336   231.332   142.481   19.587   52.114   101.706   157.153   153.536   79.610     NT2RP2000336   231.332   142.481   190.587   52.114   101.706   157.153   153.536   79.610     NT2RP2000341   37.765   23.882   31.577   11.745   14.488   23.860   24.285   20.279     NT2RP2000341   97.169   86.021   59.155   47.116   24.169   74.619   64.790   62.555     NT2RP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NT2RP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NT2RP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NT2RP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NT2RP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065		NT 2RP2000283								
NT2RP2000296										
NTZRP2000310   29.075   14.696   16.125   5.503   10.245   16.627   19.121   11.456										
## NT2RP2000310										
NTZRPZ000327										
NTZRPZ000328   36.600   35.521   50.933   15.515   23.798   33.981   22.925   32.863	40									
NTZRP2000329										
NTZRP2000333   33.894   26.367   89.382   12.302   13.127   27.377   10.155   15.517     NTZRP2000337   14.768   17.723   21.972   6.203   6.291   12.498   8.041   5.971     NYZRP2000346   53.051   82.391   46.420   15.624   13.030   26.358   27.011   31.395     NTZRP2000357   30.149   22.042   28.730   11.084   7.733   16.593   11.667   6.892     NTZRP2000358   16.228   10.853   14.700   2.291   4.114   11.789   8.150   7.184     NTZRP2000366   82.288   25.117   44.596   10.329   16.344   44.774   37.686   6.229     NTZRP2000369   21.429   15.884   19.746   6.532   11.361   9.148   7.691   12.275     NTZRP2000394   31.766   23.882   31.577   17.745   14.448   23.860   24.285   20.279     NTZRP2000396   231.332   142.481   190.587   52.114   101.706   157.153   153.536   79.610     NTZRP2000412   67.028   66.250   119.740   21.685   25.253   30.952   32.657   39.766     NYZRP2000414   97.169   86.021   59.155   47.116   24.169   74.619   64.790   62.555     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.139   27.658   7.585   9.872   17.817   19.531   15.065     NTZRP2000420   34.977   33.1										
NTZRP2000337										
NYZRP2000346   53.051   82.391   46.420   15.624   13.030   26.358   27.011   31.395		NT2RP2000333	33.894	26.367	89.382	12.302	13.127			
NTZRPZ000358 16.228 10.853 14.700 2.291 4.114 11.789 8.150 7.844 NTZRPZ000368 82.288 25.117 44.596 10.329 16.344 44.774 37.686 6.290 HTZRPZ000369 21.429 15.884 19.746 6.532 11.361 9.148 7.691 12.275 NTZRPZ000376 205.303 111.496 120.655 34.558 45.976 138.158 139.412 79.987 NTZRPZ000389 31.766 23.882 31.577 11.745 14.448 23.860 24.285 20.279 NTZRPZ000398 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610 NTZRPZ000316 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766 NTZRPZ000414 97.169 86.021 59.155 47.116 24.169 74.619 64.790 62.555 NTZRPZ000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065										
NTZRPZ000358 16. 228 10. 853 14. 700 2. 291 4. 114 11. 789 8. 150 7. 184 NTZRPZ000366 82. 288 25. 117 44. 596 10. 329 16. 344 44. 774 37. 686 6. 290 NYZRPZ000369 21. 429 15. 884 19. 746 6. 532 11. 361 9. 148 7. 691 12. 275 NTZRPZ000376 205. 303 111. 496 120. 655 34. 558 45. 976 138. 158 139. 412 79. 987 NYZRPZ000334 31. 766 23. 882 31. 577 11. 745 14. 448 23. 860 24. 285 20. 279 NYZRPZ000396 231. 332 142. 481 190. 587 52. 114 101. 706 157. 153 153. 536 79. 610 NTZRPZ000312 67. 028 66. 250 119. 740 21. 685 25. 253 30. 952 32. 657 39. 766 NYZRPZ000314 97. 169 86. 021 59. 155 47. 116 24. 169 74. 619 64. 790 62. 555 NTZRPZ000420 34. 977 33. 139 27. 658 7. 585 9. 872 17. 817 19. 531 15. 065	45									
NT2RP2000366   82. 288   25. 117   44. 596   10. 329   16. 344   44. 774   37. 686   6. 290     NT2RP2000369   21. 429   15. 884   19. 746   6. 532   11. 361   9. 148   7. 691   12. 275     NT2RP2000376   205. 303   111. 496   120. 655   34. 558   45. 976   138. 158   139. 412   79. 987     NT2RP2000394   31. 766   23. 882   31. 577   11. 745   14. 448   23. 860   24. 285   20. 279     NT2RP2000395   231. 332   142. 481   190. 587   52. 114   101. 706   157. 153   153. 536   79. 610     NT2RP2000412   67. 028   66. 250   119. 740   21. 685   25. 253   30. 952   32. 657   39. 766     NT2RP2000414   97. 169   86. 021   59. 155   47. 116   24. 169   74. 619   64. 790   62. 555     NT2RP2000420   34. 977   33. 139   27. 658   7. 585   9. 872   17. 817   19. 531   15. 065										
ETZRP200369 21.429 15.884 19.746 6.532 11.361 9.148 7.691 12.275  NTZRP2000376 205.303 111.496 120.655 34.558 45.976 138.158 139.412 79.987  NYZRP2000394 31.766 23.882 31.577 11.745 14.448 23.860 24.285 20.279  NYZRP2000396 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610  NTZRP2000412 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766  NYZRP2000414 97.169 86.021 59.155 47.116 24.169 74.619 64.790 62.555  NYZRP2000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065										
50 NT2RP2000376 205.303 111.496 120.655 34.558 45.976 138.158 139.412 79.987 NT2RP2000394 31.766 23.882 31.577 11.745 14.448 23.860 24.285 20.279 NY2RP2000396 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610 NT2RP2000412 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766 NY2RP2000414 97.169 86.021 59.155 47.116 24.169 74.619 64.790 62.555 NT2RP2000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065										
50 NT2RP2000394 31.766 23.882 31.577 11.745 14.448 23.860 24.285 20.279 NT2RP2000396 231.332 142.481 190.587 52.114 101.706 157.153 153.536 79.610 NT2RP2000312 67.028 66.250 119.740 21.685 25.253 30.952 32.657 39.766 NY2RP2000414 97.169 86.021 59.155 47.116 24.169 74.619 64.790 62.555 NT2RP2000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065				1						
50   NT2RP2000336   231, 332   142, 481   190, 587   52, 114   101, 706   157, 153   153, 536   79, 610     NT2RP2000312   67, 028   66, 250   119, 740   21, 685   25, 253   30, 952   32, 657   39, 766     NY2RP2000414   97, 169   86, 021   59, 155   47, 116   24, 169   74, 619   64, 790   62, 555     NT2RP2000420   34, 977   33, 139   27, 658   7, 585   9, 872   17, 817   19, 531   15, 065										
NT2RP2000396 231, 332 142, 481 190, 587 52, 114 101, 706 157, 153 153, 536 75, 610 NT2RP2000412 67, 028 66, 250 119, 740 21, 685 25, 253 30, 952 32, 657 39, 766 NT2RP2000414 97, 169 86, 021 59, 155 47, 116 24, 169 74, 619 64, 790 62, 555 NT2RP2000420 34, 977 33, 139 27, 658 7, 585 9, 872 17, 817 19, 531 15, 065	50									
NYZRPZ000414 97, 169 86.021 59.155 47.116 24.169 74.619 64.790 62.555 NYZRPZ000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065										
NT2RP2000420 34.977 33.139 27.658 7.585 9.872 17.817 19.531 15.065										
M12K1 2000-20 34: 311 00: 100 21:000										
NTZRP2000422   17.225   26.571   24.546   8.167   6.449   11.697   14.485   17.945										
		MT2RP2000422	17. 226	26.571	Z4. 546	8, 167	6.449	11.097	14.485	17.343

Table 82

	And I		01 1	00 750 1	51.358	87.480	90,640	100.150
		117.810	111. 501	29,759				
NT2RP2000428	56.117	63.709	38. 237	12.835	20.360	38.761	41.161	42.507
NT2RP2000438	54.621	34.534	49. 392	15.765	13.700	27.527	31.816	22.667
NT2RP2000447	41.157	17.807	23. 084	6.863	12.794	25, 289	17,738	11.474
		27.807	28. 584	7. 787	12.459	20.751	18.164	12.208
NT2RP2000448	26.410							
NT2RP2000459	44. 499	36.093	89.605	12.882	14. 284	17. 465	15. 331	9. 860
NT2RP2000479	21, 922	30. 183	53. 808	9, 553	8.835	9.648	8.854	6.739
NT2RP2000498	97. 221	94. 691	207.697	30.335	41.292	29, 900	25.090	43, 440
					10. 196	9. 766	10.763	12,056
NT2RP2000503	15.067	15.551	20.810	5. 166				
NT2RP2000510	8. 340	5. 361	8.647	4. 438	7.160	4. 784	7.812	3, 890
NT2RP2000514	10.423	8.148	14.693	2.596	1.773	12.792	6.695	2. 902
NT2RP2000516	24. 587	13.672	21.344	7.854	6, 333	13.895	7, 396	10.960
					8.071	0.000	6.857	1.961
NT2RP2000523	10.281	2.981	4. 878	1.371				
NT2RP2000533	26. 452	20.054	30. 481	4. 391	7.628	16.125	48.840	17. 396
NT2RP2000540	52. 523	22.512	28. 503	13.567	14.612	28.427	22.545	11. 372
NT2RP2000547	22.542	17.741	11.176	7.337	25,779	12, 216	8.288	6, 918
				30. 438	30.047	43.813	31,490	9.367
NT2RP2000557	91.024	63.951	163.497					
NT2RP2000558	53.959	47.359	125. 971	27.348	16.844	24. 191	17.114	21.905
NT2RP2000564	30.446	23.045,	22. 258	13.084	14. 165	16.265	14.861	11, 150
NT2RP2000565	12. 593	5.857	10. 293	5.077	0.000	4.189	5,009	9.707
				29. 211	14. 291	50. 282	32.844	34, 467
NT2RP2000583	92. 921	56.070	58. 992					2.073
NT2RP2000591	14.655	9. 331	13. 087	3.504	0.000	10.526	4. 362	
NT2RP2000599	8.002	4, 780	- 7. 951	1.807	1.614	6.232	2. 299	8. 293
NT2RP2000601	63.609	21.655	47. 106	9.673	13.430	48.855	32.575	8. 428
		37, 142	48. 248	16, 412	25. 194	51.543	39. 363	20. 157
MT2RP2000603	101.578				23. 134		30.624	31.736
NT2RP2000610	78. 342	66.011	110.636	42.146	27.855	28. 332		
NT2RP2000614	139, 380	106.590	188.604	171.750	58.678	83.079	86. 298	185. 276
HT2RP2000616	124. 143	34.073	58.053	15.031	27, 800	81.174	49.504	27.143
NT2RP2000617		37.802	37.086	17, 602	12.086	34.751	20. 157	16.389
	50.724					19,775	10. 261	13.251
NT2RP2000523	39. 247	19,740	34. 797	9.070	10. 223			
HT2RP2000634	29. 431	24.224	35. 865	13.077	19.480	16. 373	23.806	11.338
NT2RP2000536	39. 598	28.832	34. 563	11.868	13.914	14, 342	6.334	10.485
NT2RP2000638	43.027	34. 379	58. 259	14.094	15. 200	22.724	21.525	6.843
					35. 466	29. 256	23.666	11.793
NT2RP2000644	87.622	66. 336	227. 352	37.298				
NT2RP2000649	28. 849	24.035	32. 562	15. 166	18.629	25.012	15. 485	15. 528
NT2RP2000652	39, 595	25.065	30. 965	10.579	14. 587	24.849	13.667	10.824
NT2RP2000656	12.851	14.986	7. 925	2.952	4. 388	9. 997	3. 990	6.959
			7. 563	1, 152	3. 535	5, 669	3.050	2,703
NT2RP2000658	8. 192	5, 499					13. 553	39, 917
NT2RP2000663	38. 633	21.653	37. 840	5.964	12. 174	20.777		
NY2RP2000664	102.627	41.981	90.611	25. 300	30.038	73.440	66. 686	30.392
NY2RP2000668	41.209	35.434	46. 568	16.251	14.705	25. 339	29.016	11.020
		2.096	21.949	0.402	5.899	0.262	1.098	1.48
NT2RP2000678	6.908					24. 523	19.843	12.31
NT2RP2000694	47. 376	19.986	45.832	2.636	16.192			
NT2RP2000704	159, 158	114.202	205. 746	44.471	48.627	68, 161	47.919	40.34
NT2RP2000710	33, 138	26.994	21.890	10.683	6.833	17. 938	13. 596	8.070
NT2RP2000712	15.016	11.689	29. 736	12.471	8, 668	17, 629	19.970	23.79
					17.051	26.042	17. 159	21.32
NT2RP2000715	61.771	35. 912	115.757	20.470			23. 895	26.12
NY2RP2000720	38.951	26.992	43.620	14.647	11. 930	21.500		
NT2RP2000731	8.039	11.373	11.261	2.986	4.755	2. 127	4. 657	5.82
NT2RP2000739	83.662	28, 893	61.699	15.623	21.878	30.716	28. 485	17.19
NT2RP2000748	21.953	22.377	38.996	16,815	15.564	15, 846	20, 219	20,05
				1 . 4 4.4		52.159	26.941	23.86
NT2RP2000749	45.622	49.334	65. 231	13.317	57.514			9.47
			49.681	18.768	17.058	49. 245	31.463	
NT2RP2000758	79.204	43. 258	1. 43.00.				32, 438	13.13
NT2RP2000758				10.203	16.308	36.761	1 32. 430	
NT2RP2000758 NT2RP2000764	65.396	28.914	41.243				18. 850	15.66
NT2RP2000758 NY2RP2000764 NT2RP2000766	65.396 40.275	28. 914 50. 060	41. 243 83. 340	10.610	48.180	26.506	18.850	
NT2RP2000758 NY2RP2000764 NT2RP2000766 NT2RP2000777	65.396 40.275 92.029	28. 914 50. 060 39. 471	41. 243 83. 340 41. 396	10.610 32.309	48. 180 33. 513	26. 506 94. 887	18. 850 43. 480	40.21
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786	65.396 40.275 92.029 91.676	28. 914 50. 060 39. 471 61. 265	41. 243 83. 340 41. 396 70. 189	10.610 32.309 16.798	48. 180 33. 513 30. 669	26. 506 94. 887 51. 517	18. 850 43. 480 41. 968	40.21 37.84
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786	65.396 40.275 92.029 91.676	28. 914 50. 060 39. 471 61. 265	41. 243 83. 340 41. 396 70. 189	10.610 32.309	48. 180 33. 513	26. 506 94. 887	18. 850 43. 480	40.21 37.84
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786 NY2RP2000793	65.396 40.275 92.029 91.676 245.992	28. 914 50. 060 39. 471 61. 265 91. 135	41. 243 83. 340 41. 396 70. 189 151. 153	10.610 32.309 16.798 57.903	48. 180 33. 513 30. 669 62. 361	26. 506 94. 887 51. 517 191. 087	18. 850 43. 480 41. 968 132. 793	40.21 37.84 68.35
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786 NT2RP2000793 NT2RP2000796	65.396 40.275 92.029 91.676 245.992 24.053	28. 914 50. 060 39. 471 61. 265 91. 135 16. 664	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693	10.610 32.309 16.798 57.903 12.773	48. 180 33. 513 30. 669 62. 361 8. 297	26. 506 94. 887 51. 517 191. 087 14. 258	18. 850 43. 480 41. 968 132. 793 11. 004	40.21 37.84 68.35
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786 NT2RP2000793	65.396 40.275 92.029 91.676 245.992 24.053 118.982	28. 914 50. 060 39. 471 61. 265 91. 135 16. 664 88. 958	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693 221. 024	10.610 32.309 16.798 57.903 12.773 42.198	48. 180 33. 513 30. 669 62. 361 8. 297 50. 535	26. 506 94. 887 51. 517 191. 087 14. 258 65. 921	18. 850 43. 480 41. 968 132. 793 11. 004 39. 243	40.21 37.84 68.35 10.44 46.53
NT2RP2000758 NT2RP2000764 NT2RP2000766 NT2RP2000777 NT2RP2000786 NT2RP2000793 NT2RP2000796	65.396 40.275 92.029 91.676 245.992 24.053 118.982	28. 914 50. 060 39. 471 61. 265 91. 135 16. 664	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693	10.610 32.309 16.798 57.903 12.773	48. 180 33. 513 30. 669 62. 361 8. 297 50. 535 9. 968	26. 506 94. 887 51. 517 191. 087 14. 258 65. 921 23. 492	18. 850 43. 480 41. 968 132. 793 11. 004 39. 243 19. 671	40.21 37.84 68.35 10.44 46.53 6.48
NT2RP2000758 NY2RP2000764 NY2RP2000766 NY2RP2000786 NY2RP2000786 NY2RP2000793 NT2RP2000796 NT2RP200099 NT2RP2000912	65.396 40.275 92.029 91.676 245.992 24.053 118.982 23.931	28. 914 50. 060 39. 471 61. 265 91. 135 16. 664 88. 958 28. 037	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693 221. 024 26. 224	10.610 32.309 16.798 57.903 12.773 42.198 15.476	48. 180 33. 513 30. 669 62. 361 8. 297 50. 535 9. 968	26. 506 94. 887 51. 517 191. 087 14. 258 65. 921	18. 850 43. 480 41. 968 132. 793 11. 004 39. 243	40.21 37.84 68.35 10.44 46.53 6.48
NT2RP2000758 NY2RP2000764 NY2RP2000766 NY2RP2000777 NY2RP2000786 NY2RP2000793 NT2RP2000796 NT2RP200099 NT2RP2000912 NY2RP2000914	65.396 40.275 92.029 91.676 245.992 24.053 118.982 23.931 9.108	28.914 50.060 39.471 61.265 91.135 16.664 88.958 28.037 7.645	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693 221. 024 26. 224 7. 698	10.610 32.309 16.798 57.903 12.773 42.198 15.476 5.179	48. 180 33. 513 30. 669 62. 361 8. 297 50. 535 9. 968 4. 196	26. 506 94. 887 51. 517 191. 087 14. 258 65. 921 23. 492 5. 655	18. 850 43. 480 41. 968 132. 793 11. 004 39. 243 19. 671 3. 821	40.21 37.84 68.35 10.44 46.53 6.48 2.23
NT2RP2000758 HT2RP2000764 HT2RP2000766 HT2RP2000777 HT2RP2000786 HT2RP2000793 HT2RP2000796 HT2RP200099 HT2RP2000912	65.396 40.275 92.029 91.676 245.992 24.053 118.982 23.931	28. 914 50. 060 39. 471 61. 265 91. 135 16. 664 88. 958 28. 037	41. 243 83. 340 41. 396 70. 189 151. 153 26. 693 221. 024 26. 224	10.610 32.309 16.798 57.903 12.773 42.198 15.476	48. 180 33. 513 30. 669 62. 361 8. 297 50. 535 9. 968 4. 196 4. 975	26. 506 94. 887 51. 517 191. 087 14. 258 65. 921 23. 492	18. 850 43. 480 41. 968 132. 793 11. 004 39. 243 19. 671 3. 821	15.66 40.21 37.84 68.35 10.44 46.53 6.48 2.23 17.12

Table 83

	LUZABOAAAAA I	10 021	14 100 1	22 106 1	4, 470	4. 973	11.664	8. 535	5. 164
	NT2RP2000819	18. 931	14.180	22. 186					21. 582
	NT2RP2000841	28. 455	24.097	27.497	8. 335	10.021	20.722	20. 951	
5	NT2RP2000842	34. 381	17.071	34.845	8, 688	13.092	22. 498	16. 807	14. 291
3	NT2RP2000845	168, 513	153. 241	289.355	60. 191	54. 194	71.809	58. 470	61.375
	NT2RP2000863	43, 408	19.456	21.479	5. 334	8. 450	25. 326	17.757	8. 334
	NY2RP2000880	57.370	45, 920	51.291	29.897	15, 173	32.007	24. 723	27. 411
						4. 302	10.360	11. 152	10. 295
	NT2RP2000892	10.063	13.581	18, 264	3.215				12. 407
	NT2RP2000894	64. 414	18, 305	26.241	9. 579	7. 189	24. 935	24. 193	
40	NT2RP2000903	38. 945	14.595	23.755	3. 839	8.850	15.625	14. 467	11. 189
10	NT2RP2000906	43.895	24.347	34. 459	12.388	12.400	29. 304	18. 688	23. 184
	NT2RP2000910	76.036	47.430	175, 193	28. 258	21.020	30. 976	28.638	37. 229
	NT2RP2000931	68.351	104, 907	108.794	52.697	65.250	51.718	29. 344	55. 383
					6. 448	13.290	15. 553	11.313	11, 145
	HT2RP2000932	30.706	39,023	31.030			32.600	18.046	21. 528
	NT2RP2000938	55.079	37.641	47.798	12.045	19.899			
	NT2RP2000943	64.610	32.689	54. 181	11.802	18.241	33.817	55. 424	23. 572
15	NT2RP2000957	20, 426	12.332	17. 780	3, 161	5.343	6.479	8.015	4. 052
	NT2RP2000958	74. 825	23.934	37.910	10.227	22.164	41.633	29. 369	21. 256
	NT2RP2000959	15. 840	25.063	17.980	5. 521	4.208	9. 176	3. 539	6. 349
	NT2RP2000965	52. 687	40. 458	51.330	27.882	16.372	29. 535	32. 993	35. 643
						36.529	42.914	24. 489	33. 313
	NY2RP2000970	84. 866	72.715	196.279	29. 249			20.702	18.360
	MY2RP2000973	42.690	30.786	42.102	8.964	13.498	23. 369		
20	NT2RP2000985	33. 281	22.399	26, 930	8.628	4.869	20.022	22.445	14.030
	NT2RP2000987	47.736	66. 487	94. 477	25. 911	19.844	27. 890	23.633	33. 551
	NT2RP2000997	42.801	43.070	56.966	15. 270	16. 292	49.613	53. 625	99. 622
	NT2RP2001024	47.605	28.976	34, 658	13.810	14, 526	32.054	39. 269	22.962
	NT2RP2001028	32. 502	24,770	88. 599	12. 437	11.259	13.181	13.919	9. 824
	NT2RP2001036	206. 163	234.625	568. 339	116.746	85.893	125. 996	88. 623	100.568
0.5				31, 356	6.335	15. 429	17.827	107.341	12.412
25	HT2RP2001039	26.909	37.527				31. 422	25. 682	20. 463
	NT2RP2001044	51. 134	33.868	42.988	9.015	23.633			44. 980
	NT2RP2001056	84. 875	95.778	164.256	33. 325	35.039	45. 764	30.831	
	NT2RP2001065	57. 092	61.052	49.599	18.558	20. 229	29. 013	30.628	32. 966
	MT2RP2001067	17. 223	18.596	14.258	5. 284	6,021	4. 582	10.045	8.782
	NT2RP2001070	92.615	68.975	230.584	37.646	41.225	36. 295	43. 293	26.959
20	NT2RP2001081	134.654	80, 124	269.700	35. 425	37.697	42.849	35. 852	45. 105
30	NT2RP2001087	54. 476	40.059	74.079	12.377	21.043	25.654	22.663	15. 956
	HY2RP2001094	11.558	8. 400	11.506	4.416	3, 583	4. 503	4. 258	5. 446
					36.523	37.388	40.013	41.672	39. 968
	NT2RP2001119	66. 924	57.741	177.347			16.757	28.865	13.483
	NT2RP2001127	52. 585	39, 380	36.247	9.959	18.625			
	NT2RP2001133	94.638	97, 465	155. 477	25. 417	36, 346	28.836	28. 731	38. 218
35	NT2RP2001137	61.770	53.486	51.726	12.991	40.072	20. 107	24. 686	30.341
33	NT2RP2001142	54. 131	38.507	34.342	8. 552	14.688	17. 434	23.807	20.602
	NT2RP2001149	96.617	49. 914	71.348	17.462	14.077	23.064	30. 576	22.427
	NY2RP2001168	313.055	217.008	205.763	65. 294	77, 914	146.883	169. 121	159. 484
	NT2RP2001173	25. 149	27.272	22.710	16.143	12.538	13. 238	14. 902	9.473
		21. 134	17.440	22.879	11.089	14. 190	18. 125	50.600	22.839
	NT2RP2001174			84.254	29. 471	35. 438	70. 558	65. 859	57. 928
40	NY2RP2001184	99.803	60.549			9,060	20. 695	9. 289	15, 340
	NT2RP2001196	19.492	14.580	26.749	5.551			33. 102	31.874
	NT2RP2001200	39. 331	44. 223	52.647	14.745	26.231	26.146		
	NT2RP2001218	32.396	16.531	28.960	21.387	13.855_	8.618	18.872	11.236
	NT2RP2001223	86.393	27. 183	45.400	14. 290	23.545	53.375	28.096	26.084
	NT2RP2001225	223.868	143.880	155.700	46.575	60.808	148.876	100.150	92.898
	NT2RP2001227	100.969	51.807	65.094	19.398	24. 302	57.877	36. 375	36.204
45	NT2RP2001232	49.733	30. 526	64, 154	11.691	29.542	27. 238	22. 294	35. 950
			36. 288	152.784	58. 935	18.921	38.027	28. 582	69.539
	NT2RP2001233	42.734				8. 746	38. 272	8. 565	38.035
	NT2RP2001245	28. 251	16. 266	32.594	18.419			31.550	37. 123
	NT2RP2001246	24.708	44. 426	35,600	19, 345	16.443	35.994		
	NY2RP2001268	44. 328	34. 570	58. 263	12.894	20.636	54.014	31.715	54.645
	NT2RP2001270	37.478	15. 214.	29.740	12.749	67.060	24.740	26.469	21.423
50	NT2RP2001276	15. 931	7. 906	12.674	9. 235	4.389	12.549	17. 273	10. 235
	ST2RP2001277	22. 937	21, 147	33.688	12.970	7.618	12.672	2.878	16.107
			20. 688	27.890	13. 340	37.970	38, 444		34.029
	NYZRPZ001290	66.867				12.979	24. 206	7.863	10. 592
	HT2RP2001295	22.777	21.635	31.845	7.387				399.018
	NT2RP2001297	105.753	198.744	183. 982	210.648	32.481	152.615		
	HT2RP2001301	47.099	37. 782	53.504	25. 117	15. 392	49. 389	38.668	29. 281
66									

· Table 84

			136 000	204 (14	05 020	120 170 1	315 760 7	000 070 I	146 642
	NT2RP2001312	493.097	175. 989	324.513	96.070	132.150	315.768	282. 270	146. 542
	HT2RP2001327	188.839	50.032	95.732	33.162	58.029	112.666	87.335	71.442
5	NT2RP2001328	177.255	162. 267	495.438	96. 591	104.203	93.675	57.120	68.709
	NY2RP2001341	196.358	92. 246	40.237	32.288	34.069	91.368	77.221	45.741
	NT2RP2001347	148, 143	157. 594	486.643	72.828	67.867	81.012	36.464	72. 260
	NY2RP2001366	160. 323	170. 553	496.412	116.205	96.521	146.562	77.918	108.669
	MT2RP2001378	217. 791	51.524	110.978	31.128	51.690	147. 191	118.132	56. 442
	MT2RP2001381	16, 578	13.963	19.068	15.119	9. 576	8. 483	2.703	10.418
10	NT2RP2001388	84.013	52.476	228. 213	47.276	49.007	52.881	33.168	44. 592
	NT2RP2001391	806.136	1438. 949		960. 225		1160.112		1127.811
	NT2RP2001392	56. 943	65. 258	70. 204	19.962	26.883	46.456	23. 261	14. 231
	NT2RP2001394	104. 258	120.852	350.764	78.963	59.635	75. 686	42.505	53.751
	NT2RP2001397	37.759	22.378	38.780	40. 524	15.364	21.089	16.393 11.796	16.560 22.055
	NT2RP2001400	24. 214	10. 586	19.685	10.414	12. 173	24.380 29.278	20.555	45,713
15	NT2RP2001408	34. 405	28. 262	69.823	33.071	21.313	33. 427	30.009	41.019
	NT2RP2001420	74.700	70.462	212.932	44. 495	49.469	21.739	12.751	10. 452
	NT2RP2001423	20.045	17. 202 91. 272	38.815 206.946	16.204 51.057	11.082 36.829	49.854	34. 587	57.012
	NT2RP2001427	88.620	45. 465	55. 112	19.580	15. 421	24. 651	10.915	29. 985
	NY2RP2001428 NT2RP2001436	47.617 19.654	25. 606	50. 345	11. 202	18.548	32.033	22.720	5. 351
	NT2RP2001440	11.871	12. 123	19.145	7.724	5.414	7.413	19.955	16.145
20	NT2RP2001445	11.934	7.217	22.053	5. 885	6.872	7.794	3.377	20.818
	MY2RP2001449	20. 271	20.423	53.385	13.242	8.026	-8. 439	6.342	9.186
	NT2RP2001450	47.497	32, 496	58. 237	18.660	21.208	28.880	23.620	30.207
	NT2RP2001467	40.279	40.050	115.089	25. 502	21.744	18.716	21.445	37.772
	NY2RP2001469	66.890	35.784	93. 465	23. 588	33.470	54.095	54. 103	33. 386
	NY2RP2001480	69.698	53.669	54,777	16.208	26.373	44. 943	30.622	26.208
25	NT2RP2001495	14, 156	12.199	18.018	8.178	14.762	10.694	8.800	14.613
	NY2RP2001499	40.983	50.266	57.334	23. 302	22.298	37.271	26.788	35, 187
	NT2RP2001506	83.528	66.377	104. 162	41.795	65.692	61.567	55.661	35.667
	CT/2RP2001508	25.746	36.879	44.112	33, 620	14.149	23.999	19.783	36.174
	NT2RP2001511	231.898	147.751	199.611	46. 927	77. 381	122.787	130.829	108.021
	MY2RP2001514	121.671	47. 391	103.398	24, 149	31.957	72.965	63. 365	38. 173
30	NT2RP2001520	38.773	20.470	34.140	14. 159	13.166	19.602	22.077	7.741
	NY2RP2001526	102.469	96.418	139.331	62.159	83.922	85.309	60. 450	66.763
	NT2RP2001529	189. 308	69.082	103.704	31.713	54. 543	173.158	96.700	74. 482 17. 343
	MT2RP2001536	22.047	14. 186	19. 269	9.553	7.196	16.531	13.646 133.592	422. 435
	MT2RP2001538	123, 315	222.563 33.999	281.173 42.028	191.775	90.257	199. 255 31. 438	29. 406	24. 085
	MT2RP2001547	45.201 146.079	68.501	131.623	35, 625	46.061	88.704	90.584	78. 703
35	NT2RP2001562	53.975	35, 141	47. 262	23. 297	18.361	43.041	30.635	47.577
	NY2RP2001566	55, 453	48.563	91.463	37. 157	27, 507	54.780	37.595	42. 563
	NY2RP2001569	131.940	142.523	361.640	62, 136	60.136	90.021	46.500	62.567
	NT2RP2001576	103.537	76.306	58. 434	23.607	34.646	91.306	67. 270	45. 219
	DT2RP2001581	149.528	208.681	239.575	139, 522	72.883	196.577	126.583	231.505
	NT2RP2001597	52.409	27.790	43.630	13.807	18.650	35.875	23.646	43.012
40	CT 2RP2001601	33.796	37.430	70.562	17.535	15.251	22. 525	13.760	29.828
	NT2RP2001613	10.438	5. 350	6.715	3. 155	6.423	9.119	9.830	14.501
•	NT2RP2001628	87. 399	43.401	48.713	17.774	25.577	50.117	31.175	117.652
	NT2RP2001634	38.792	56.546	47.793	23. 992	16.006	30.530	21.235	42.849
	NY2RP2001635	63.818	69.842	156. 279	31.411	35.011	40.036	38.853	22.210
	HT2RP2001660	31.554	25.538	25. 905	6.081	11.137	20.048	20.365	48. 159 43. 557
45	NT2RP2001862	122.557	88.914	242. 932	52.514	43.761	63.759	56.518 20.869	25. 608
	MY2RP2001653	33.056	34. 206	58.783	11.163	16.477	39.485 33.037	25.666	35. 948
	NT2RP2001672	51.656	46.965	140.882	31.231	26.225	3. 149	4, 861	6.461
	NT2RP2001675	8.589	6.791	12.510	1.982	5.806 27.233	40.323	46. 466	47.741
	NT2RP2001677	70.100	49.851	68.423 193.110	17.674 58.566	46.915	65.668	62.835	64. 186
	NT2RP2001678		86.779	25.445		10.015	9. 959	25.390	9. 277
50	NT2RP2001683	16.088	14.728 54.743	185.463		33, 217	64. 457	41.391	50. 672
	NT2RP2001699 NT2RP2001707	94.748	66.728	100.874		34. 234	58.720	45. 599	68. 302
	NT2RP2001720	81.079	33.745	39.415		16. 907	38.973	31.931	30. 227
	ST2RP2001721	73.164	35. 354	62.124		28.378	69.464	66. 522	35. 468
	NT2RP2001740	23.081	30. 430	27. 131	12.949	12. 248	21.055	20.053	28.546
	THIEN COULING	1 60.001	1 55. 750	1 27.191	1,5.373	1.2.2.70			•

Table 85

	COTATE AND A STATE OF	164 378 1	- PT PAA T	121 YES 1	45 COO T	10 124 1	06 212 1	62 222 1	25 004
	NT2RP2001748	164.370	51.538	151.756	22.608	36.134	86.312	52. 323	35.004
	NT2RP2001755	10.363	5.707	7. 354	3.303	2.490	20. 122	2.133	9. 371
•	NT2RP2001762	10.743	10.704	7. 130	4.777	5.648	16.360	7.429	3.763
5	NT2RP2001768	122.047	71.860	129.000	29.098	38.722	67.999	58. 129	48.111
	NT2RP2001769	29. 307	28.706	32. 455	11.608	15.175	19, 399	20.505	29. 469
	NT2RP2001784	18. 824	19. 122	24, 434	8. 167	13.814	14.835	14, 266	10, 332
	NT2RP2001805	111,510	63.886	82. 038	33.170	41.704	47, 921	62.218	54, 508
	NT2RP2001813	15.000	10.225	13. 797	4. 221	9. 786	3, 548	11.805	8.246
			12.403	19. 383	6.848	8. 320	6. 884	10.608	15. 163
10	NT2RP2001817	14.005					12.858	13.055	9, 397
	NY2RP2001818	30.494	21.374	23. 441	6.808	14. 438			
	NT2RP2001837	153. 478	143. 980	348, 522	65.249	55, 344	69.434	48. 042	62.813
	NY2RP2001839	68. 237	44.005	65. 237	21.186	23.824	37.874	35. 524	54. 235
	NT2RP2001861	45. 604	33. 558	72.763	21.180	25. 185	40. 479	31. 542	29. 326
	NT2RP2001869	79. 101	52.967	123. 399	29.766	25.811	40.870	38.251	38. 150
	NT2RP2001876	20, 847	28, 536	35. 991	18.044	13, 257	29. 195	20.056	35.651
15	NT2RP2001878	105. 429	34.989	86.887	21.675	33. 547	76.806	64.301	35, 521
	NT2RP2001881	25, 562	5. 186	16, 935	8.594	6.002	8.017	5.474	16.018
	NT2RP2001883	162. 487	96.494	76. 800	26.663	40.257	93,069	57.806	50, 662
				18, 175	19.127	0.000	34.665	13. 313	27.989
	NT2RP2001884	40.027	29. 435			9.345	26. 427	24.717	30. 448
	NT2RP2001885	41.527	29. 494	60, 284	13.719				
	NT2RP2001898	152.071	65.585	135. 420	33.617	41.173	112.042	64. 105	57.703
20	NT2RP2001900	20.075	16. 336	54. 207	10.431	9.790	20.098	19. 168	30. 123
	NT2RP2001903	389. 922	207.168	314.475	131.627	170.618	361.733	261.185	289.339
	HT2RP2001907	118. 240	77.557	213,664	50.816	46. 691	58.895	52.711	56.061
	NT2RP2001915	29. 335	9. 240	29. 213	5. 804	10.101	8.718	14.671	15.535
	NT2RP2001921	70.657	42.199	23. 786	27.411	23.817	52.083	27.655	30.244
	NT2RP2001926	86.771	11.953	10.434	11.123	10.945	27.144	37.077	26.703
25	NY2RP2001933	210. 457	80.003	159. 875	38.312	53. 192	114.539	90.251	48.849
	NY2RP2001936	9, 271	13.789	9.841	9.560	6.311	8.706	3.968	4.244
	NT2RP2001943	129.800	151.136	357.167	96. 135	99, 997	227.342	186.800	161.131
	NT2RP2001946	36. 700	27.839	38. 317	18.830	11.786	20.082	32.636	29.552
	NT2RP2001947		40. 322	58. 260	17. 399	25, 524	30.411	31. 309	15, 258
		49. 825			5.855	16.449	8.590	3, 943	39. 227
	NT2RP2001948	6.858	5. 149	39. 338		55.776	144.746	109.645	45. 142
30	NT2RP2001956	204, 499	97.036	150. 184	34.215				
	NT2RP2001969	63.044	42.091	64.895	18. 446	22.555	64. 128	29.876	27.818
	NT2RP2001976	8.014	10.925	13. 322	14.259	2.776	2.729	6.432	21. 452
	NT2RP2001978	60.910	40. 459	87.051	23.282	28.689	25. 497	33. 528	35. 507
	NT2RP2001985	73.126	35.661	72.052	21.029	30.385	52.485	45. 885	41.899
	NT2RP2001991	32.897	34.028	33.239	10.548	15.586	20.531	18. 489	33. 157
	NY2RP2001997	38. 265	33.006	69.711	20.057	29.835	29.074	30. 213	39. 156
35	NT2RP2002015	341,660	572.382	454. 288	330.114	80. 297	356.270	346.254	476.966
	NT2RP2002017	33, 468	25.736	55. 897	13. 982	18.424	23.720	12.540	17.897
	NY2RP2002025	201.899	111.493	125. 922	38.775	57.018	118.130	92.718	55.437
	NT2RP2002030	147. 806	150.643	447. 960	95.773	104. 163	95. 260	65.007	88.254
	NY2RP2002032	170. 695	55.335	101.868	30. 495	58.859	127.664	86.380	56.817
				481. 152	84.872	61. 493	72.667	37.144	74. 278
40	NT2RP2002033	147.111	92.379	17. 284	5.762	7. 552	5. 398	10.885	30. 538
· •		15.097		25. 228	11.030	8. 158	11.642	14. 255	15. 385
	NT2RP2002046	15.094	19. 275					10. 225	22. 164
	MT2RP2002047	19. 261	15.499	12.076	6.530	14. 384	9.918		
	NT2RP2002050	71. 226	75.633	97.017	33. 238	36. 421	49.003	42.580	45.656
	NT2RP2002052	75.004	67.588	69.616	25. 123	25. 691	49. 820	32.819	35. 546
	MT2RP2002058	9.803	11, 955	11.648	6.527	5. 940	8. 570	15.678	16.434
45	NT2RP2002080	147. 927	40.191	79.254	17.661	30.022	83.968	55. 933	35.933
	HT2RP2002063	8.334	10.615	17. 124	3.910	9.032	6. 499	6.095	43.967
	MT2RP2002066	85, 296	31.968	71.727	16.697	28. 928	52. 589	40.814	37. 383
	NT2RP2002070	24.791	21.309	66.961	13.511	11.537	11.893	11.300	28.065
	NY2RP2002076	28. 441	16.541	17.729	6.137	10.519	13.321	11,910	9. 273
	NT2RP2002078	75. 992	38.941	77. 227	23. 502	30.063	65.434	39.358	28, 599
60						11.345	7, 129	16.510	27. 362
50	NT2RP2002079	15. 378	6.595	12.418	5.815				
	NT2RP2002099	78. 520	17.490	39, 514	8.705	17.165	51.830	37.473	36.146
	NT2RP2002105	45. 619	26.109	41.837	15. 263	18. 979	33.970	43.561	26. 203
	NT2RP2002115	4. 270	4. 361	2.711	1.795	2.838	1.055	1.725	0.659
	NT2RP2002124	9. 528	14. 188	19. 276	6.091	6. 494	4. 045	5. 259	20. 125
	HT2RP2002137	42. 205	16.239	58.339	7. 326	12.132	22.097	14.684	13.003
55									

## Table 86

	NT2RP2002139	134. 906	45, 566	87.430	23. 245	34.053	84. 189	66.692	39. 692
	NT2RP2002154	73.877	40, 956	58.084	17. 359	21.276	53. 977	32.847	25. 981
-	NT2RP2002155	312.813	448. 404		246. 578	165. 102	220. 200	117.089	396.951
5	NT2RP2002172	30. 233	30.674.	.55. 659	15. 991	11.479	50. 028	14. 567	76. 933
							23. 455	23.637	18. 868
	NT2RP2002185	35. 127	22.047	31.771	9.436	13.476			78. 046
	NT2RP2002188	281.595	70. 032	141.729	44. 333	58.463	164. 587	118.094	
	NT2RP2002192	28.830	19. 474	84. 849	18. 536	13. 421	8. 267	10.908	25. 727
	HY2RP2002193	51. 545	23. 270	33.672	10.534	17.989	33.897	31.972	33.050
10	NT28P2002208	28.592	23. 922	46.625	15. 986	13.078	25. 948	18.689	40. 263
	NT2RP2002219	13. 529	18. 299	23.304	8. 697	7.005	20. 832	6. 994	4. 421
	NT2RP2002231	3. 623	9. 145	18.238	6.451	5. 394	5. 290	2.444	1.640
	NT2RP2002232	41.922	30.600	40.665	10.290	12.646	31.637	16.070	23. 193
	NT2RP2002235	25. 174	12.829	11.461	1.747	8.624	10. 246	12.594	16.053
	NT2RP2002239	123.883	99.627	183.537	54. 220	35, 311	68. 845	72.486	114. 538
15	NY2RP2002252	173. 209	45. 051	80. 502	16. 296	33.546	82. 843	82.445	52. 048
	MT2RP2002256	6.776	3. 892	12.301	3.488	7.236	6.566	9. 391	- 9. 526
	HY2RP2002257	14.914	18.059	11.330	3. 304	7.442	11.747	12.965	136.057
	NT2RP2002259	25.623	20. 902	41.590	9. 164	7.958	18.892	22.893_	29. 020
	NT2RP2002264	35.467	21.380	27.456	3.962	7.884	26.448	8. 234	20. 246
	NT2RP2002267	99.224	90. 968	353.970	55. 091	43.373	63.895	30.469	55. 401
20	HT2RP2002270	12.038	20. 146	13.141	7.551	3. 523	7.777	6.701	19. 108
20	MY2RP2002281	49.615	38, 410	43.936	21.926	17. 935	51. 455	14.825	35. 239
	MT2RP2002288	18.840	15. 310	15. 237	4.623	6.951	4. 505	6.438	4. 321
	NT2RP2002292	70.138	79.487	98.062	32.152	32.815	48. 306	41.287	55. 682
	NT2RP2002299	28.411	21.790	28. 450	15.762	10.016	23.812	12.394	31.923
	NY2RP2002304	17.776	27. 505	25. 401	9.478	10.570	14. 112	10.173	10. 213
	NY2RP2002312	32.053	25.004	19.733	5.118	10.392	41.845	21.011	16.815
25	NT2RP2002316	15.618	29, 406	20.363	11.321	29.588	16.866	17.862	43.519
	NT2RP2002325	32.321	23.882	28. 597	6.692	9.875	26. 435	21.261	36.989
	NY2RP2002333	117.384	75.765	92.724	37.475	55. 245	56.768	79.089	134. 509
	ST2RP2002371	35.025	49.789	54.117	20.073	31.179	10, 486	24. 281	48. 279
	MT2RP2002373	73.024	55.638	58.797	24.729	33.686	48.754	58. 440	58. 483
	NT28P2002381	4.610	6.610	5. 950	2.906	4.109	10.398	7.035	3. 142
30	NT2RP2002385	73.600	28.798	39, 973	10.268	23.738	57.377	29.062	18. 367
	NT2RP2002394	4.749	3.341	5. 573	1.941	3.227	11.225	3.017	2.611
	NY2RP2002408	30.199	16.610	24.803	8.840	17.966	22.778	22.751	14.463
	NT2RP2002409	466.226	415.995	746.844	183.086	221.410	247.550	216.812	235.852
	NT2RP2002424	73.955	40.022	38.701	11.417	27.269	38.757	36. 192	25.977
	NT2RP2002426	42. 246	46.209	138.641	18.951	43.167	21.993	14.146	29.925
35	NT2RP2002429	38.798	37.515	37.290	13.976	31.959	40. 592	16.576	28. 408
	NY2RP2002437	41. 182	44, 109	103.486	16.002	6.706	22.769	11.006	18. 502
	NT2RP2002439	300.787	110.081	147.018	33.619	60.331	171.025	155. 332	90. 923
	NT2RP2002442	51.674	59. 162	57.683	24. 271	21.412	43. 427	38. 136	78. 512
	NT2RP2002457	87.804	91.782	200. 265	. 53. 883	50. 903	42.083	43.069	58. 125
	NT2RP2002464	97.665	38.612	69.981	20.743	31.183	66.794	48.779	34.847
40	MY28P2002475	87.229	49. 226	48.473	16. 952	38. 579	51.432	45.816	27.604
40	NT2RP2002479	43.495	20.334	24. 184	10. 295	13.868	35. 366	19. 292	22.684
	MY2RP2002487	95.041	44. 922	72.897	21.815	31.046	43.590	37. 943	47.177
	NT2RP2002498	32.022	15.599	33.143	12.736	8.092	15. 582	24.301	15. 152
	NT2RP2002503	143. 137	80.337	119.421	48. 392	35.509	96.570	63.743	69.363
	HY2RP2002504	28.779	12.130	143.283	15.019	25.676	16.936	24.798	15.731
45	NT2RP2002510	389.826	185.539	464.842	123. 573	125.657	192.079	171.751	115.972
45	NY2RP2002520	28.465	20.629	47.388	22.909	14.948	38.504	25.659	37. 802
	NT2RP2002527	82.404	66.911	163.583	35.753	34.220	51.754	33.562	45. 539
	NT2RP2002533	453.205	209.788	357.064	113, 267	150.283	251. 157	252.839	188.717
	NY2RP2002537	39.475	40.266	89.504	25.635	20.657	31.517	13.708	23. 210
	HY2RP2002542	68.000	79.669	80.611	82.297	29.448	38.068	33.806	62.834
	NT2RP2002546	27.656	17.241	60, 211	11.584	0.000	26.089	6. 935	5. 274
50	HY2RP2002549	41.394	22.287	57.825	30.309	7.713	40.681	12.786	23.580
	NT2RP2002564	135.808	83.403	115.471	41.607	30.969	95. 939	62.575	50. 150
	NT2RP2002591	34. 917	38.054	103.943	37.411	25. 346	30.888	24, 127	41.780
	NY2RP2002595	29. 155	28.991	47, 139	17, 440	18.504	24.511	28.272	25. 178
	NT2RP2002602	62. 164	42.498	49. 596	18. 894	40.679	48.767		7.981
	NT2RP2002606	23. 368	18.641	18.058	7, 405	14. 392	5.066	8. 402	
55	MIENT LUVE BUG	1 63.300	1 10.041	1 ,5. 555	1 7, 403	1	1		
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Table 87

NT2RP2002609	51.566	22.622	50.513	17.534	20.249	18.692	26.812	44. 49
T2RP2002618	54.802	20.530	64. 541	20. 957	20. 177	31.222	20.834	32.08
T2RP2002621	108.854	151,631	361.642	75.866	73.104	87.556	37.662	72. 94
NT2RP2002643	79, 459	49.749	159.326	32.265	31.588	30.054	50.389	48. 13
NT2RP2002672	97. 309	70.875	124.816	41.317	54.912	65. 362	54. 912	61. 19
		27.357			16. 225	18. 592	18.872	41.66
NT2RP2002673	33, 731		31.454	11.741				
NT2RP2002674	13.503	12.059	23. 980	5. 008	15. 903	5. 926	8.720	8.88
NT2RP2002686	45.156	22.604	57.057	22. 253	22. 373	30.389	27.672	13.37
NT2RP2002688	85. 273	71.163	154.737	61.783	35. 115	56. 421	42.460	68. 11
NT2RP2002595	80.865	40.613	62.941	16.213	22.197	43. 453	30, 540	28. 17
NT2RP2002701	68.274	58.034	54, 220	24, 008	29.811	75. 585	54.744	29.99
NT2RP2002706	66.710	49.408	147.083	42.409	25.501	40.462	31, 482	31.67
NT2RP2002710	876.030	389.806		246.642	312.053	990.051	876.290	401.33
NT2RP2002711	120. 344	48.897	112.902	26.906	37.076	81.599	62, 600	40.80
					10.968	18. 932	17.127	19.19
NY2RP2002727	19. 985	16.809	28.658	5.885				
NY2RP2002734	84. 484	81.389	244. 997	57.973	45. 229	35. 711	33.199	39.65
NT2RP2002736	18.170	7.757	29.873	5. 264	10.456	10. 179	9. 257	11.01
NY2RP2002740	13.219	14. 424	23.343	12.863	6.975	8. 152	8.795	7.77
NY2RP2002741	77.823	67.266,	223.592	33.955	36.594	51.261	45. 295	14.04
NT2RP2002750	140.558	111.369	512,500	99.367	68, 412	72.711	76.999	72.28
NT2RP2002752	177.349	105, 312	290. 520	63, 592	64, 508	103.376	92. 228	65.84
NT2RP2002753	131.824	60.851	110.980	32.981	43.667	85.850	102, 908	117.42
NY2RP2002760	130.675	58, 967-	119.405	28. 837	37.588	59. 420	51, 267	51.70
			32.873		12.332	10.357	15. 988	25.04
NT2RP2002769	19.077	14.018		14. 190				77.04
NT2RP2002778	38.616	37.548	30.303	18. 271	16.022	71.865	31.460	
NT2RP2002791	95.319	55. 458	105.096	34, 190	38.076	66.995	54. 639	45.5
NT2RP2002800	90.052	59, 554	197.798	40.413	37. 123	87.119	52.880	48.1
NT2RP2002805	14.997	12.041	9. 573	4.470	8.397	5. 324	5. 699	14.60
NT2RP2002811	84. 563	36.955	70.308	17.273	24.509	89.018	46.163	49.1
NT2RP2002824	44. 392	48, 364	75, 269	21.980	25.621	56.385	42.073	38.1
NT2RP2002839	45. 683	28, 499	42.893	12.083	18.567	22.078	23.650	21.6
NT2RP2002845	46. 337	22.545	45, 003	11.450	16.060	6.978	26, 900	14.5
NT2RP2002857	26.773	11.114	27.648	7. 358	7.968	15.413	17.314	11.9
	122. 430	114, 903	392.000	81.893	61,001	82.758	60.301	50. 3
NT2RP2002862							14.768	24.8
NT2RP2002880	46.913	32.677	29.822	12.750	16.704	35. 359		
NT2RP2002885	24. 335	26.185	27.174	10.146	19.062	54. 580	55. 170	22.5
HT2RP2002891	33.411	27.772	38.018	14.600	16. 632	38.658	34, 150	26.2
NY2RP2002907	31.117	36, 465	35. 948	13.227	13.010	49. 335	37. 225	26.7
NT2RP2002925	30.213	17.281	33.298	11.072	11.726	25. 559	24.754	17.4
NT2RP2002927	21.224	35.383	40. 539	21, 437	7.365	35. 485	14.771	39.4
NT2RP2002928	13.771	14, 521	49, 574	11.977	6.869	9. 129	7, 289	8.0
NT2RP2002929	21.741	22.530	32.027	7.934	12.601	20. 143	13, 573	25. 5
NT2RP2002934	63. 248	35, 331	42.688	10.849	16.987	39, 637	27.937	23.4
		30.833	62.082	15, 330	19.313	35. 512	35.749	25. 2
NT2RP2002939	53.914				53.315	49, 000	38. 922	90. 3
NT2RP2002942	82.129	82.694	187.805	50. 572				26.6
NT2RP2002954	33. 490	25. 335	35.779	11.591	11.217	27. 293	16.672	
NT2RP2002959	18.029	22.305	18. 230	8.391	14.540	12.392	9. 227	31.2
NT2RP2002974	34.775	17.807	29.755	6.220	18.382	28. 562	36.888	41.1
NT2RP2002976	7. 266	6.893	13.152	2.886	5. 205	17.007	6.657	17.8
NT2RP2002979	156, 906	139, 229	395. 529	82.939	71.144	104. 220	76.074	81.3
NT2RP2002980	98.467	79.422	285. 396	49. 557	40.675	57.510	33.004	50.4
MY2RP2002986	210. 452	66.962	105.842	25. 570	34.404	156.863	99. 482	35. 9
NT2RP2002987	170.131	130.848	355. 987	114.067	85.014	125. 562	105. 241	119.4
NT2RP2002988		33.804	42.437	7.516	22.093	78. 216	26.257	53. 4
	35. 092 41, 408				13.951	19. 869	17.068	17.7
		20.150	29.978	8. 083				61.9
NT2RP2002993			265.303	52.674	45. 922	52. 225	38. 486	
NT2RP2002993	91.683	72.701				22.796	23.430	31.
NT2RP2002993 NT2RP2003000 NT2RP2003000	91.683 19.429	42, 300	26.458	14.959	11.323			1 7A '
NT2RP2002993 NT2RP2003000 NT2RP2003000 NT2RP2003020	91.683			14.959	198. 298	95. 120	89. 298	
NT2RP2002993 NT2RP2003000 NT2RP2003000	91. 683 19. 429 146. 283	42. 300 83. 102	26.458 231.026				89. 298 24. 460	
NT2RP2003000 NT2RP2003000 NT2RP2003000 NT2RP2003020 NT2RP2003032	91. 683 19. 429 146. 283 42. 858	42, 300 83, 102 35, 052	26. 458 231. 026 46. 187	31. 287 15. 872	198. 298	95. 120	89. 298	29.
NT2RP2002993 NT2RP2003000 NT2RP2003008 NT2RP2003020 NT2RP2003032 NT2RP2003032 NT2RP2003034	91. 683 19. 429 146. 283 42. 858 97. 685	42, 300 83, 102 35, 052 100, 455	26.458 231.026 46.187 302.158	31. 287 15. 872 45. 216	198. 298 16. 376 40. 853	95, 120 25, 572 44, 346	89. 298 24. 460 20. 833	29. 6 60.
NT2RP2002993 NT2RP2003000 NT2RP2003000 NT2RP2003020 NT2RP2003032 NT2RP2003034 NT2RP2003034	91. 683 19. 429 146. 283 42. 858 97. 685 32. 097	42, 300 83, 102 35, 052 100, 455 30, 146	26.458 231.026 46.187 302.158 30.859	31. 287 15. 872 45. 216 9. 131	198. 298 16. 376 40. 853 14. 406	95. 120 25. 572 44. 346 14. 312	89. 298 24. 460 20. 833 25. 483	29.6 60. 23.
MT2RP2002993 NT2RP2003000 NT2RP2003000 NT2RP2003020 HT2RP2003034 NT2RP2003034 NT2RP2003042 NT2RP2003050	91.683 19.429 146.283 42.858 97.685 32.097 43.965	42, 300 83, 102 35, 052 100, 455 30, 146 23, 480	26. 458 231. 026 46. 187 302. 158 30. 859 42. 356	31. 287 15. 872 45. 216 9. 131 12. 150	198. 298 16. 376 40. 853 14. 406 15. 913	95. 120 25. 572 44. 346 14. 312 20. 938	89. 298 24. 460 20. 833 25. 483 29. 611	29.6 60.3 23.8 20.9
NT2RP2002993 NT2RP2003000 NT2RP2003000 NT2RP2003020 NT2RP2003032 NT2RP2003034 NT2RP2003034	91. 683 19. 429 146. 283 42. 858 97. 685 32. 097	42, 300 83, 102 35, 052 100, 455 30, 146	26.458 231.026 46.187 302.158 30.859	31. 287 15. 872 45. 216 9. 131	198. 298 16. 376 40. 853 14. 406	95. 120 25. 572 44. 346 14. 312	89. 298 24. 460 20. 833 25. 483	74.3 29.6 60.3 23.8 20.9

Table 88

	NT2RP2003099	60.090	61 064	107 021	20 002	70 400	69 700 1	30 145 1	46.759
		69. 980	61. 964	197.831	28. 962	29. 485	52.756	36.145	46.753
	NT2RP2003108	22.037	23. 450	29.734	12.784	12. 243	25. 414	19. 582	14. 441
5	NT2RP2003115	175. 202	76.490	219.003	26.090	53. 025	89. 403	96.086	53. 165
	NT2RP2003117	132. 572	135. 106	428, 449	65.631	66.802	77.649	41. 504	75. 169
	NT2RP2003121	77. 521	49.860	42.009	15.143	26. 745	31.652	32.041	27. 916
	NT2RP2003125	35. 377	29.656	27. 135	9, 957	16. 383	12.805	20. 265	8. 252
	NT2RP2003127	29. 566	16.867	20. 397	5. 212	10, 531	18. 240	19. 752	7. 540
	NT2RP2003129	50.461	54. 112	157.477	25.025	29.892	16.686	23. 103	33.770
10	NT2RP2003137	8.001	18.759	14, 140	10.321	7.469	15. 281	5. 429	3. 225
70	NT2RP2003138	52. 296	44. 278	85. 267	21.446	22.368	30.612	24. 709	34. 031
	MT2RP2003146	55. 329	37.398	52.403	14. 492	12.222	29.608	23, 329	32.663
	NT2RP2003148	150. 386	104. 523	330.270	60. 524	70. 523	90.836	76. 602	100. 291
	NT2RP2003150	26. 432	11. 157	23, 761	15.678	11.132	36.468	7. 133	18, 954
	NT2RP2003157	58. 172	46.518	64. 963	42. 288	23. 422	50.314	42. 129	48. 145
45	NY2RP2003158	44. 248	20. 906	37.740	8. 136	17, 954	27. 119	19.062	38. 471
15	NT 2RP2003161	19. 274	11.968	16.062	2.701	7.578	17.086	7.441	31.024
	NT2RP2003164	49. 401	19.110	28.830	12. 219	12.819	22. 155	19.787	34.090
	NT2RP2003165	89. 985	65. 955	218. 487	37.132	35, 205	34. 406	24. 887	33, 303
	NT2RP2003177	43. 596	22.142,	51. 196	11.148	3.934	15. 303	13.349	69.154
	NT2RP2003179	69.718	46.328	169, 618	30.883	22.456	37.444	43. 967	45.776
•	NT2RP2003194	144. 137	17. 980	22. 293	13.420	10.852	20. 144	19.065	43.611
20	MT2RP2003206	7.840	5. 369	10.850	6.014	4. 029	11, 290	7.725	3.709
	NT2RP2003210	51. 322	21.586	38. 521	12.974	17.884	37.608	30. 477	29.805
	HT2RP2003227	42. 906	18.716	24. 162	17. 143	9, 513	37.425	15. 949	23, 165
	NT2RP2003228	58. 612	29.572	62.903	22. 926	28. 577	30.449	37. 367	63. 378
	NT2RP2003230	5. 885	10. 431	148. 181	5. 253	9. 252	9.617	6. 228	22. 492
	NT2RP2003231	69. 197	41.691	59, 459	34.789	15. 272	58.827	33.617	37.859
25	NT2RP2003237	30. 563	38.860	123, 572	28.832	11.050	15. 189	9. 580	23.097
	NT2RP2003239	33. 469	21.053	50.845	20. 348	11.513	25.692	7.484	35. 924
	NT2RP2003243	145, 467	34. 182	76.360	17.705	28.702	66. 482	55. 093	28. 921
•	MT2RP2003255	29. 516	23. 976	32.673	9.710	15.918	17.608	20. 157	14. 165
	NT2RP2003267	65. 087	29.515	67. 969	24. 282	21.518	34.797	27. 241	43.679
	NT2RP2003272	41.457	22.351	19.055	27.076	19.762	28.028	26. 982	45. 977
30		107. 913	82.634	92. 986	31.633	32.424	67.812	26. 460	53.116
	NY2RP2003280 NT2RP2003286	19. 151 21. 848	14.918	20.689	11.633	7. 567	43. 338	5. 070	12.961
			17.740	29. 829	11.104	6. 965	28.110	26.734	26. 233
	MT2RP2003293	94. 719	83. 407 16. 886	364. 260	76.134	56. 105	78.539	44. 376	97.047
	HT2RP2003297	17.874		18.717	18. 256	19.625	15.088	25. 617 10. 162	16. 166
	MT2RP2003300	9. 592 15. 144	10.816	15. 547 26. 519	2. 211 10. 354	5. 615 14. 045	8. 461 6. 847	8.974	5.662 11.058
35	NY2RP2003302	22. 071	15. 550	64. 230	26.397	10. 289	12.880	11.722	68. 523
	NT2RP2003307	22. 086	9.418	17. 120	5. 220	6. 112	15.691	17. 396	7. 096
	NY2RP2003308	17. 436	24. 315	20. 930	11.886	7.814	20. 422	12.860	31.766
	1372RP2003311	22.001	9. 144	13.842	5. 360	10.074	18.616	5. 176	21. 146
	NT2RP2003329	44. 872	14.471	19.961	10.976	13.401	22.292	12.093	14.770
	MT2RP2003339	20. 422	19.625	85.412	16. 458	12.443	17.818	9, 125	13, 152
40	MT2RP2003345	23. 118	8. 297	17.237	4. 695	8. 379	12.952	12. 259	23. 215
	MY2RP2003347	12. 389	4.636	9. 822	7.720	7. 500	12.461	7. 182	16.011
	MT2RP2003367	10.794	19.368	21.160	7.884	14. 120	12, 142	14.419	13, 409
	NT28P2003369	41. 141	18. 327	38, 318	11.072	14, 356	33.971	28. 126	19.613
	HT2RP2003383	55. 891	32.218	76,058	21.558	27.536	76.861	50. 564	36. 175
	NT2RP2003390	73.620	57.765	91.034	41.124	35. 539	63.744	46.234	42.766
45	MT2RP2003391	241.564	161.239	277.051	75. 828	95. 432	220.668	152.546	143, 981
	NT 2RP2003393	11.758	13.507	20.112	4. 687	11.809	12.940	19.991	21.749
	MY2RP2003394	7. 323	9.816	9.506	2.871	10.713	1.307	6. 346	14.753
	NY2RP2003401	25. 259	3.938	8.376	2.832	4. 096	7.246	16. 169	7.442
	NT2RP2003403	31.239	26, 205	109.072	18.680	14.206	9. 380	14. 946	8.745
	NT2RP2003433	79.603	33.408	70.460	19.431	29. 526	42.730	34.783	28.629
50	MT2RP2003445	38. 525	33.248	95, 090	23.648	21.333	27.951	21. 347	33. 662
- •	NT2RP2003446	67. 228	39.971	49. 302	18.878	21.829	54. 339	39.113	29.464
	NT2RP2003456	1. 902	13.833	10.178	7.437	1.522	5.049	1.410	3. 486
	HT2RP2003466	72.001	27.022	47.862	12.506	26.814	66.543	51.004	41.515
	NY2RP2003469	35, 915	29. 791	90.766	19. 568	17.254	24.857	16. 952	39. 575
	MT2RP2003470	20. 820	31.916	84.744	64.680	20.126	61.522	22. 215	98. 657
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55									

Table 89

										10 010
		NT2RP2003471	7.424	5. 547	6.488	7.037	5.447	6.505	7.782	10.212
		NT2RP2003480	78.094	65, 408	137.798	31, 787	40.594	58.633	37.776	39.678
			15. 982	11.924	14, 233	7.870	5.725	11.076	8. 329	14, 404
5		NT2RP2003495								
		NT2RP2003499	55. 449	13.382	25. 597	4. 229	14. 517	54. 430	36.252	15. 105
		NT2RP2003505	55. 425	27.024	46.996	11.964	7.933	31.002	31.997	27. 989
		NT2RP2003506	29.029	19.815	26.696	9.949	12.205	23. 185	12, 152	24. 906
		NIZKPZUUJSUO								
		NT2RP2003511	85. 237	37.479	50. 383	22.212	25. 152	50.854	41.079	36.551
		NT2RP2003513	2, 085	4. 521	4. 122	3.531	5.027	3.740	2.918	7.377
		NY2RP2003517	37. 834	17. 587	35. 502	11, 597	12.069	30.516	43.651	39.873
10										
		NT2RP2003522	24. 832	37.794	30.938	13.985	21.613	21.384	15.975	15.713
		NT2RP2003525	112.839	77.947	318.516	53.968	64.300	64.511	45. 220	44. 281
		NT2RP2003533	95. 494	87.932	267,080	44.833	35. 543	46.891	33, 401	37.402
							24. 451	41.018	38.504	56.566
		NT2RP2003541	59. 237	40. 256	51.598	18.653				
		NT2RP2003543	60. 456	24.016	25.862	11.661	16.145	17.623	31.288	25. 312
		NT2RP2003545	5. 111	9. 859	11.338	12, 197	5.950	2.774	8.060	34.030
15		NT2RP2003559	26. 905	22. 287	37.874	13.292	12.911	24.477	17, 350	31.685
		NT2RP2003564	29. 146	18.045	64.896	13.749	13.213	15.703	17.055	25.744
		NT2RP2003565	71.340	105.907	131.344	34, 826	44.614	78.728	62.826	61.650
		HY2RP2003567	70.892	54. 381,	72.715	19.440	21.968	61. 162	50.325	46.459
							4. 227	5. 271	7,753	9,628
		NT2RP2003575	8. 045	11.848	16.656	3.697				
		NT2RP2003576	94. 175	119. 128	189.789	159.528	39.210	94. 530	84.153	280.017
20		NT2RP2003579	55. 985	110.923	72.170	19.865	32.853	121.326	99.589	58.803
		NT2RP2003581	72. 231	34. 935	63.218	15. 922	25.161	44. 829	45.801	38.825
									62.139	47.892
		NT2RP2003587	109. 102	46, 403	76.235	20. 483	28.667	127. 344		
		HT2RP2003590	27.361	26.330	26.653	9.837	5.016	24. 313	17.397	36. 147
		NT2RP2003593	98.848	66.189	91, 401	17.565	31.030	61.583	54, 982	56.233
		HT2RP2003596	20. 156	17.830	46.567	15. 376	7.364	8.849	10.462	35. 925
							33.831	64, 394	76.259	72.122
25		MY2RP2003599	99. 163	72.506	53.708	30. 551				
		NT2RP2003600	39.566	25. 200	27. 397	13.373	16.019	22.567	30. 947	25. 783
		NT2RP2003604	30, 188	48.497	24.769	15.941	13.513	20.832	18.908	35.739
		NT2RP2003629	12.593		13.520	5.134	7. 235	8.896	12.558	21.197
				10.012			24.270		32. 166	28. 383
		NT2RP2003630	55.769	31.553	55.456	13.290		37.506		
		NT2RP2003643	20. 532	14.638	38.212	9.363	17.760	18.713	18. 506	19.629
20		MT2RP2003655	46.795	29.612	38. 397	10.145	18.688	20. 220	24. 997	18.685
30					21.831	11.981	11,047	39. 022	14.701	15.715
		NT28P2003664	23.372	28.188						
		NT2RP2003668	98.074	77.678	215.011	48.838	58.733	45. 358	46.022	49.968
		MT2RP2003687	36.469	27.937	30.101	11.600	12.659	14.676	15.349	16, 155
		MT2RP2003691	57.166	66.814	140. 266	28.579	24.877	10.915	18.651	30.704
						37.740	29.269	33. 935	36.174	35, 262
		MT2RP2003702	77.231	74.259	157.835		29.209			
35		MT2RP2003704	33.958	19. 273	90. 406	13.087	15.614	12. 526	13. 208	27.631
33		HY 2RP2003706	15.581	9.802	10.782	1.905	1.888	20.850	8.045	6.106
		NT2RP2003713	16.960	13. 155	19.058	12.333	6.597	11.248	12.533	12.834
							25. 935	21. 990	15.804	26.140
		NT2RP2003714	58. 106	48, 190	156.974	28. 216				
		NT2RP2003727	16.878	30.048	13, 471	24.840	10.360	26. 581	2.051	18.209
		HY2RP2003737	35.097	27.626	24.696	15.279	8.490	48. 230	26.577	18.778
		HT2RP2003751	24. 927	12.926	14. 285	5.654	5. 352	15.115	11.036	11.385
40						31.937	11.912	70.013	35.412	\$0.086
. •		MT2RP2003760	61.964	14.851	34.689					
		NT2RP2003764	70.923	28.030	49.140	23. 190	33. 253	31.845	28.042	21.978
		HT2RP2003789	42.617	20.886	27. 599	7.054	10.396	11.852	16.178	10.912
		NT2RP2003770	137.506	66. 296	82. 283	29.001	19,657	59. 586	43.465	55.063
						21.542	23.944	31.481	38. 443	30,003
		NT2RP2003777	79.392	37.432	49.453					14 558
		MY2RP2003781	113.598	78.822	248.846	43.005	41.064	65. 158	51.558	43.936
45		NT2RP2003785	39.008	38.895	81.842	23.800	81.398	60.210	21.078	32.965
		NT2RP2003793	29.403	32.842	38.373	11.279	11.070	27.094	13, 519	16.114
	•							54.142	52.055	74.576
		NY2RP2003806	141.377	86.683	300. 547	56.391	57.427			
		NT2RP2003825	200.861	142.661	421.147	81.431	83.143	96.953	65.464	115.589
		NT 2RP2003840	100.905	61.436	80.952	27, 801	38.812	73.708	55.685	43.672
						48.707	32.982	109.107	66.696	63, 138
		NT2RP2003857	135.915	99.087	88.444				39.367	23.246
50		NT2RP2003859	112.898	91.670	144.716	35. 434	18.445	66.240		
		MT2RP2003871	16, 891	14.873	18.946	20.075	9.742	10.433	6.276	13. 332
		NY2RP2003876	20.553	18.667	33, 132	17.736	9.744	22.067	11.629	10.917
						7. 186	11.534	5. 285	2.003	13.835
		NT2RP2003878	10.935	24. 440	15,728					
		NY28P2003885	86.851	91.093	40.636	9.621	12.995	23. 247	25.798	7. 129
		NT2RP2003898	42.684	30. 561	43.471	13.576	37.187	19.007	22.509	33. 529
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#### Table 90

NT2RP2003902	147.643	124, 985	109, 475	45. 984	48.594	124.353	51.962	58. 344
NT2RP2003912	125. 311	242, 124	511.945	129. 243	109. 998	129.880	47.537	95. 222
NT2RP2003931	26.887	8.179	6, 459	2.307	5, 260	8. 153	1.858	3. 142
NT2RP2003940	186. 397	64,618	262.034	55.607	30.649	41.635	23. 343	65.087
NT2RP2003950	36. 158	19, 195	49, 413	13.592	20.939	19.343	26.770	21.989
NT2RP2003952	15. 955	17.931	35, 750	13.974	12.406	27. 300	20.083	13.016
NT2RP2003968	45. 877	22.833	13. 459	11.361	12.355	12.353	12.010	25, 113
NT2RP2003976	37, 958	44, 808	95. 495	38.986	28.544	21.209	8. 325	15. 117
MT2RP2003981	38.654	43.006	57, 657	15. 338	29. 345	30.659	23.563	25.867
NT2RP2003984	132.353	65.644	60.516	16.394	44. 914	84.097	45.289	33.280
NT2RP2003986	186.062	146,313	421. 324	109.891	71.468	70.656	43.927	53, 945
NT2RP2003988	112.131	82.329	348. 163	81.784	60.909	64. 387	44. 174	58.384
NT2RP2004013	35. 821	31.054	41.104	24.447	20.809	33.899	21, 394	38.113
NT2RP2004014	51.068	77.076	125. 407	38. 647	29. 948	34.055	26.943	33.783
NT2RP2004036	34. 592	12.491	12.862	9.166	7.965	9, 771	12.722	18. 319
NT2RP2004041	61.828	31.728	66. 443	16.578	28.668	39.049	31.113	30. 197
NT2RP2004042	95. 416	34.628	56. 458	18. 193	31.581	50.180	28.757	19.510
NT2RP2004049	30.836	31.163	33.858	10.780	19. 423	28.518	29. 763	8. 339
NT2RP2004060	33. 939	22.080	47.086	13.117	10.598	29.819	24. 922	24. 074 25. 483
NT2RP2004066	36. 939	51.977	61.500	23. 281	20.470	26.729	15. 403 30. 550	18. 563
NT2RP2004069	29. 217	33.889	47. 332	22.168	14.676	23.715 5.069	6.316	20. 634
NT2RP2004076	9. 020	12.153	35. 232	4. 198 4. 309	9. 970 8. 489	27.512	· 5. 327	10. 188
NT2RP2004080	23.022	8.835	21.995 83.806	31.063	33. 602	10.431	18. 338	56.090
NT2RP2004081	38.786 47.764	30.091	36. 354	14.003	22.548	26. 497	22.648	13. 621
NT2RP2004108	28.744	38. 559	67.714	34. 947	23.442	39.884	20.636	48. 103
NT2RP2004108	43.031	24.659	37.232	12.008	12. 194	23. 487	10, 186	21.361
NT2RP2004130	62.738	36.522	73.772	37.407	24. 390	44.094	20.478	34. 479
ST2RP2004133	163.939	56.278	112.008	40.808	61.092	157. 167	95. 384	52. 343
MT2RP2004141	49.570	22.611	50.916	9.793	20.924	53. 203	22.033	30. 465
NT2RP2004142	34.850	23.492	33.078	17, 102	15. 132	27.703	11.237	17. 601
NT2RP2004152	14. 256	11.207	21.943	19.655	8.860	14.997	12.981	8. 353
NT2RP2004165	147.447	92.813	238. 228	40.497	54. 357	70.413	30.081	44. 940
HT2RP2004170	107.111	64.978	194.673	41.028	56.020	66.291	58.470	56. 553
NT2RP2004172	22.440	1.5. 213	19.562	6.795	12.099	15. 400	14.334	12.024
NY2RP2004176	120, 902	23.723	54.734	12.552	24. 966	70.512	39.664	28. 280
NT2RP2004179	72.406	30. 327	45. 178	12.821	11.733	33.905	35. 842	30.011
NT2RP2004187	25, 235	21.870	33.704	11.364	19.908	8.982	12.208	16.442 36.649
NT2RP2004190	33.406	32.037	37.882	8. 251	10.063	16.897	16.826 59.372	84. 014
MT2RP2004194	84.064	81.541	54.017	35. 398	25. 386	70.700 83.939	40.164	46. 168
NT2RP2004196	105.711	65.320	61. 236	35. 178	35. 795 46. 886	102.336	55. 538	55. 936
NT2RP2004205	144. 445	71.761	300. 198 14. 703	38.897 6.333	7.074	34. 908	17.403	14. 550
NT2RP2004207	34.894 63.802	12.571 26.160	69.559	17, 665	24. 160	72.242	27. 469	21.672
M12RP2004232	19.053	14.404	25.695	7.555	9.877	15. 593	12.523	32.679
NT2RP2004232	49.739	30.594	47.640	22.915	18. 596	31, 416	32.672	84. 520
NT2RP2004240	43. 946	56.977	36.742	39. 556	38. 450	39.881	22.758	41.302
HT2RP2004242	24.272	10.675	24, 496	11.743	14.023	31.038	18.900	15. 124
NT2RP2004245	18.673	23.813	15.945	12.936	16.016	18. 326	7. 178	10.903
HT2RP2004270	234, 182	227.894	511.563	104.046	110.474	124. 225	90.436	89. 248
NT2RP2004300	59, 573	43,407	77.768	15.466	13. 124	34.892	25.094	19. 570
NT2RP2004304	30.539	31.035	68.652	13.187	14.829	18. 430	12.663	17.214
NT2RP2004313	52.639	26.629	35.836	12. 439	13.307	42.833	29.621	25.693
NT2RP2004315	7.937	6.053	8. 996	2.798			1.817	5.009 6.754
MT2RP2004321	16.873	18.267	25. 584				12.417	17, 712
NT2RP2004336	27.640	16.775	31.426	5.804			18.808 70.845	110.855
NT2RP2004339	253.896	255.780					11.378	12. 471
MY2RP2004347	39, 311	42.402			14. 095		22. 223	23.600
NT2RP2004364	71.148	60.019					18.665	18. 356
NT2RP2004365	27.548	25. 940					15, 740	27. 122
HT2RP2004366	34.341	34.055				34, 007	21.569	14.963
NT2RP2004373	28. 456	29.195						28.153
NT2RP2004375	22. 258	23.633						16.693
NT2RP2004389	26, 163	1 41.010	11.340	111.240	1 .0.001			